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(54) Title: MYCOBACTERIUM TUBERCULOSIS GENES ENCODING PROTEIN ANTIGENS

(57) Abstract

Mycobacterium tuberculosis genes encoding five immunologically relevant proteins have been isolated by systematically screening a lambda gt11 recombinant DNA expression library with a collection of murine monoclonal antibodies directed against protein antigens of this pathogen. One of the *M. tuberculosis* antigens, a 65kD protein, has been shown to have determinants common to *M. tuberculosis* and *M. leprae*. In addition, genes encoding proteins of other mycobacteria (*M. africanum*, *M. smegmatis*, *M. bovis* BCG and *M. avium*) have been isolated. Isolation and characterization of genes encoding major protein antigens of *M. tuberculosis* make it possible to develop reagents useful in the diagnosis, prevention and treatment of tuberculosis. They can be used, for example, in the development of skin tests, serodiagnostic tests and vaccines specific for tuberculosis.

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MYCOBACTERIUM TUBERCULOSIS GENES AND
ENCODING PROTEIN ANTIGENS

Description

Background

05 Tuberculosis was the major cause of infectious mortality in Europe and the United States in the 19th and early 20th centuries. Dubos, R. and J. Dubos, The White Plague: Tuberculosis, Man and Society, Little Brown & Co., Boston, MA, (1952).
10 Today, it remains a significant global health problem.

15 For example, in the United States there are over 20,000 new cases of tuberculosis diagnosed annually. In addition, the steadily declining incidence of tuberculosis evident in preceding years appears to have changed course, reaching a plateau in 1985 and showing an increase in the first half of 1986. Centers for Disease Control, Morbidity/Mortality, Weekly Report, 34:774 (1986); and Centers for Disease Control, Morbidity/Mortality, Weekly Report, 35:774 (1986).

20 Worldwide, tuberculosis remains widespread and constitutes a health problem of major proportions, particularly in developing countries. The World Health Organization estimates that there are ten million new cases of active tuberculosis per year and an annual mortality of approximately three

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million. Joint International Union Against Tuberculosis and World Health Organization Study Group, Tubercle, 63:157-169 (1982).

05 Tuberculosis is caused by Mycobacterium (M.) tuberculosis or Mycobacterium (M.) bovis, which are the 'tubercle bacilli' of the family Mycobacteriaceae. M. bovis is a species which causes tuberculosis in cattle and is transmissible to humans and other animals, in whom it causes tuberculosis. At 10 present, nearly all tuberculosis is caused by respiratory infection with M. tuberculosis. Infection may be asymptomatic in some, but in other individuals, it produces pulmonary lesions which lead to severe debilitation or death. Resistance to 15 tuberculosis is provided by cell-mediated immune mechanisms.

20 Mycobacteria are aerobic, acid-fast, non-spore-forming, non-motile bacilli with high lipid contents and slow generation times. M. leprae is the etiologic agent of leprosy and, among the other mycobacteria, the only major pathogen. Bloom, B.R. and T. Godal, Review of Infectious Diseases, 5:765-780 (1983). However, other mycobacterial species are capable of causing disease. Wallace, R.J. et.al., 25 Review of Infectious Diseases, 5:657-679 (1984). M. avium, for example, causes tuberculosis in fowl and in other birds. Members of the M. Avium-intracellularare complex have become important pathogens among individuals with acquired immunodeficiency syndrome (AIDS). Certain groups of 30

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individuals with AIDS have a markedly increased incidence of tuberculosis-as well. Pitchenik, A.E. et. al., Annals of Internal Medicine, 101:641-645 (1984).

05 Diagnostic and immunoprophylactic measures for mycobacterial diseases have changed little in the past half century. Tuberculin, developed by Koch as a cure for tuberculosis in the late 1800s, is an M. tuberculosis filtrate of complex and poorly-defined composition. It is used as a skin test antigen to detect prior exposure to the bacillus. Enrichment of the protein fraction of this material in the 1930's produced the purified protein derivative (PPD) which is still used to diagnose exposure to 10 tuberculosis. Seibert, F.M. et.al., American Review of Tuberculosis, 30(Suppl.):705-778 (1934). Its usefulness is limited, however, by its lack of specificity and its inability to distinguish active disease from prior sensitization by contact with M. tuberculosis or cross-sensitization to other myco-bacteria. Young, R.A. and R.W. Davis, Proceedings of the National Academy of Sciences, USA, 80:194-20 1198 (1983).

Bacille Calmette Guerin (BCG), an avirulent strain of M. bovis, has been used widely as a live vaccine against tuberculosis for over 50 years. 25 Calmette, A., C. et.al., Bulletin of the Academy of

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Medicine Paris, 91:787-796 (1924). During that time, numerous studies have shown that BCG has protective efficacy against tuberculosis. These studies are reviewed by F. Luelmo in American Review of Respiratory Diseases, 125(pt. 2):70-72 (1982). However, more recently, a major trial of BCG in India indicated that such a vaccine was not protective against tuberculosis in this setting. World Health Organization WHO Technical Report Series, 651 (1980). Presently available approaches to diagnosing, preventing and treating tuberculosis are limited in their effectiveness and must be improved if a solution is to be found for the important public health problem tuberculosis represents worldwide.

Summary of the Invention

The present invention is based on the isolation of genes encoding immunogenic protein antigens of the tubercle bacillus Mycobacterium tuberculosis (M. tuberculosis). Genes encoding such protein antigens have been isolated from a recombinant DNA expression library of M. tuberculosis DNA. Genes encoding proteins of four additional mycobacteria have also been isolated and restriction maps produced.

In particular, genes encoding five immunodominant protein antigens of the tuberculosis bacillus (i.e., those M. tuberculosis proteins of molecular weight 12,000 daltons (12kD), 14kD, 19kD, 65kD and 71kD have been isolated by probing a lambda gt11 expression library of M. tuberculosis DNA with

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monoclonal antibodies directed against M.
tuberculosis-specific antigens.

Recombinant DNA clones producing the specific
05 antigenic determinants recognized by the monoclonal
antigens were also isolated in this manner. DNA
from such recombinant lambda gt11 clones was mapped
with restriction endonucleases; the restriction maps
for genes encoding the five immunodominant protein
10 antigens (i.e., genes encoding the 12kD, 14kD, 19kD,
65kD and 71kD proteins) were deduced. The nucleo-
tide sequence of three of the genes have been deter-
mined and, in each case, the amino acid sequence of
the encoded protein has been deduced.

Brief Description of the Drawings

Figure 1 shows restriction maps of M. tuberculosis DNA. Recombinant DNA clones isolated with
15 monoclonal antibodies directed against the 12kD,
14kD, 19kD, 65kD and 71kD protein antigens were
mapped with restriction endonucleases. The insert
DNA endpoints are designated left (L) or right (R)
20 in relation to lac Z transcripts which traverse the
insert from right to left. Restriction sites are
represented as follows: A, Sal I; B, BamHI; E,
EcoRI; G, BglII; K, KpnI; P, PvuI; S, SacI; X, XbaI.
25

Figure 2 shows arrays of antigens from M.
tuberculosis recombinant DNA clones probed with
rabbit hyperimmune serum. The code of the recombi-
nant DNA clones shown on the numbers filter is: 1,
Y3275; 2, Y3274; 3, Y3279; 4, Y3277; 5, Y3247; 6,
30 Y3272; 7, Y3150; 8, Y3254; 9, Y3147; 10, Y3163; 11,

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Y3179; 12, Y3191; 13, Y3252; 14, Y3178; 15, Y3180; 16, Y3143; 17, lambda gt11. Clones 1, 5, 6, 7, 9 and 16 are M. tuberculosis recombinants described in the following section. Clones 10, 11, 14 and 15 are 05 M. leprae recombinants expressing epitopes of the 18kD, 28kD, 36kD and 65kD antigens, respectively. Clones 2, 3, 4, 8, 12, 13 are uncharacterized recombinants from the lambda gt11 M. tuberculosis and M. leprae libraries. Clone 17 is a non- 10 recombinant lambda gt11 control.

Figure 3 shows arrays of recombinant mycobacterial antigens probed with monoclonal antibodies to assess the extent of cross-reactivity between recombinant protein antigen of M. tuberculosis and 15 of M. leprae. The array of clones is identical to that shown in Figure 2. Antibody probes and the antigen sizes recognized are: 1, IT-11 (71kD); 2, IT-31 (65kD); 3, IT-16 (19kD); 4, IT-1 (14kD); 5, IT-3 (12kD).

20 Figure 4 shows restriction maps of DNA encoding four proteins (71kD, 65kD, 19kD and 14kD) of M. tuberculosis and four proteins (71kD, 65kD, 19kD and 14kD) of M. bovis BCG. Restriction sites are represented as follows: A, AatII; B, BamH1; C, 25 BcII; D, DraIII; E=EcoRI; G, BglII; H, HinfI; K, KpnI; P, PstI; S, SalI; V, PvuI and X, XhoI.

Figure 5 is a comparison of restriction maps of the gene encoding the 65kD protein of 6 mycobacteria (30 M. leprae, M. tuberculosis, M. africanum, M. bovis BCG, M. smegmatis, M. avium). Restriction sites are

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as follows: B, BamHI; K, KpnI; N, SacI; P, PvuI; S, SalI; X, XhoI.

Figure 6 is the nucleotide sequence of the region containing the M. tuberculosis 19kD gene.
05 The deduced amino acid sequence of the encoded protein is also represented (protein start position, nucleotide 1110; protein stop position, nucleotide 1586).

Figure 7 is the nucleotide sequence of the 10 region containing the M. tuberculosis 71kD gene and the deduced amino acid sequence of the encoded protein.

Figure 8 is the nucleotide sequence of the region containing the M. tuberculosis 65kD gene.
15 The deduced amino acid sequences of the two long open reading frames are presented in one letter code over (540) or under (517) the appropriate triplets.

Detailed Description of the Invention

The invention described herein is based on the 20 isolation of genes encoding immunogenic protein antigens of the bacillus M. tuberculosis, which is the major etiologic agent of tuberculosis. In particular, it is based on the isolation, using monoclonal antibodies directed against M.
25 tuberculosis-specific antigens, of genes encoding five immunogenic protein antigens of the tuberculosis bacillus; these five antigens are immunodominant. Immunogenic antigens are those which elicit a response from the immune system.
30 Immunodominant protein antigens are immunogenic

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antigens against which the immune system directs a significant portion of its response. Genes encoding M. tuberculosis antigens of molecular weight 12,000 daltons (12kD), 14kD, 19kD, 65kD and 71kD were
05 isolated in this manner.

Isolation and characterization of major protein antigens of M. tuberculosis, as described herein, make it possible to develop more effective tools for the prevention, diagnosis, and treatment of tuberculosis.
10 Identification and isolation of genes encoding five immunodominant M. tuberculosis protein antigens, as well as of the five protein antigens, are described below; uses of the genes and encoded products are also described.

15 M. bovis BCG DNA clones were also isolated for the genes encoding the 71kD, 65kD, 19kD and 14kD proteins. In order to compare M. bovis BCG and M. tuberculosis genes encoding proteins of similar molecular weight, restriction endonuclease maps were
20 determined for DNA segments containing each of the genes. Restriction maps for each of these genes is represented in Figure 4.

In addition, DNA clones were isolated for the genes encoding the 65kD protein from M. africanum,
25 M. smegmatis and M. avium. Restriction endonuclease maps were determined for DNA segments containing each of these genes. The restriction maps for these genes, as well as for the genes encoding the 65kD protein of M. tuberculosis, M. bovis BCG and M. leprae, are represented in Figure 5.

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I. Construction of a recombinant expression library of M. tuberculosis DNA

A recombinant DNA expression library of M. tuberculosis DNA was constructed using lambda gt11. 05 The library was constructed with M. tuberculosis genomic DNA fragments in such a way that all protein-coding sequences would be represented and expressed. Young, R.A., B.R. Bloom, C.M. Grosskinsky, J. Ivanyi, D. Thomas and R.W. Davis, 10 Proceedings of the National Academy of Sciences, USA, 82:2583-2587 (1985).

Lambda gt11 is a bacteriophage vector which is capable of driving the expression of foreign insert DNA with E. coli transcription and translation 15 signals. Lambda gt11 expresses the insert DNA as a fusion protein connected to the E. coli Beta-galactosidase polypeptide. This approach ensures that the foreign DNA sequence will be efficiently transcribed and translated in E. coli. This approach is also useful in addressing the problem of the highly unstable nature of most foreign proteins; 20 fusion proteins are often more resistant to proteolytic degradation than is the foreign polypeptide alone. Lambda gt11 and the E. coli strain used 25 (Y1090) have been described previously. Young, R.A. et al., Proceedings of the National Academy of Sciences, USA, 80:1194-1198 (1983); Young, R.A. and R.W. Davis, Science, 222:778-782 (1983). The teachings of these publications are incorporated 30 herein by reference. The library constructed in this manner has a titer of 1×10^{10} pfu/ml. and

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contains approximately 40% recombinants with an average insert size of 4kB.

II. Screening of the lambda gt11 M. tuberculosis library with antibody probes

05 Murine monoclonal antibodies to protein antigens of M. tuberculosis were used individually to probe the M. tuberculosis recombinant DNA library. This work is described below and with specific reference to the 65kD antigen in the Exemplification. The antibodies used as probes and the sizes of the antigens to which they bind are shown below.

		<u>M. tuberculosis</u>	
		<u>Antibody</u>	<u>Antigen</u>
	IT-3		12kD
15	IT-20		14kD
	IT-19		19kD
	IT-27		19kD
	IT-17		23kD
10	IT-29		23kD
	IT-15		38kD
15	IT-21		38kD
	IT-23		38kD
	IT-13		65kD
20	IT-31		65kD
	IT-33		65kD
25	IT-11		71kD

Engers, H.D. et al., Infectious Immunology,
51:718-720 (1986).

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All monoclonal antibodies were used at approximately 1:200 to 1:300 dilution in 50mM Tris-HCl pH8/150 mM NaCl/.05% Tween 20.

Screening of the lambda gt11 recombinant DNA library was performed as described by Young et al. 05 in Proceedings of the National Academy of Sciences, USA, 82:2583-2587 (1985), the teachings of which are incorporated herein by reference. One modification was made in the method described by Young and 10 co-workers: 1% bovine serum albumin was used in place of 20% fetal calf serum to decrease background.

Briefly, cloned lambda gt11 recombinants were arrayed on lawns of E. coli Y1090. The phage were 15 grown, antigen expression was induced and the antigens were blotted and probed with serum. Detection of signal-producing plaques was performed with a biotinylated secondary antibody system (Vectastain, Vector Laboratories, Burlingame, CA) or 20 with an alkaline phosphatase conjugated secondary antibody system (Protoblot, Promega Biotec, Madison, WI), both used according to manufacturer's instructions. Signal-producing clones were isolated using antibodies directed against protein antigens of 25 molecular weight 12kD, 14kD, 19kD and 65kD and 71kD. In each case, similar numbers of clones were 30 isolated in screens of approximately 10^5 recombinant plaques. DNA clones encoding the 23kD and 38kD antigens could not be detected with these antibodies, possibly because the native epitope is modified or topographically complex, or because the

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antigen-antibody interaction is too weak to be recognized by current detection methods.

III. Probing of Arrays of lambda gt11 DNA Clones with Antibody Probes

- 05 0.2 ml of a saturated culture of Y1090 was added to 2.5 ml of molten LB soft agar, poured onto 100 mm plates containing 1.5% LB agar and allowed to harden at room temperature for 10 min. 100 ul of phage plate stock containing approximately 10^{11} pfu/ml of the lambda gt11 DNA clones of interest were placed into alternate wells of 96-well tissue culture plates. A multi-pronged transfer device was placed briefly in the wells containing phage and then touched lightly to the surface of the plate onto which the soft agar had been poured. The plates were then incubated at 42°C for approximately 3 hours, at which point clear plaques approximately 5mm in diameter were visible. The plates were then overlayed with nitrocellulose filters saturated with 10mM isopropylthiogalactoside (IPTG) and incubated at 37°C for 3.5 hours. Subsequent processing of filters for detection of antigen was identical to the procedures described for screening of lambda gt11 library with antibody probes.
- 10 15 20 25 Immunoscreening of the lambda gt11 library to isolate clones reactive with monoclonal antibodies specific for the 65kD antigen is described in the Exemplification.

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IV. Recombinant DNA Manipulation

DNA from recombinant lambda gt11 clones was isolated and mapped with restriction endonucleases by standard techniques. Davis, R.W. et al.,
05 Advanced Bacterial Genetics: A Manual for Genetic Engineering, Cold Spring Harbor (1980).

Figure 1 shows the genomic DNA restriction map deduced for each of the genes encoding the five M. tuberculosis antigens and illustrates how each of
10 the cloned DNAs aligns with that map. All clones isolated with monoclonal antibodies directed against any single antigen align with a single genomic DNA segment. This indicates that all clones were
isolated because they express the protein of interest rather than an unrelated polypeptide containing
15 a similar or identical epitope. In addition, this result suggests that each antigen is the product of a single gene.

The orientation of each DNA insert in the
20 recombinant clones was determined by restriction analysis. Only among the clones for the 65kD antigen were the inserts found in both possible orientations relative to the direction of lac z transcription in lambda gt11. This suggests that
25 this protein can be expressed in E. coli from signals independent of those provided by lac z. Similar results have been obtained for recombinant DNA clones encoding the 65kD antigens of M. bovis and M. leprae. Thole, J.E.R. et al., Infectious Immunology, 50:800-806 (1985); Young, R.A. et al.,
30 Nature, 316:450-452 (1985).

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The nucleotide sequences of three regions of the M. tuberculosis DNA were determined: 1) the region containing the M. tuberculosis 19kD gene; 2) the region containing the M. tuberculosis 71kD gene; 05 and 3) the region containing the 65kD gene. The three sequences are represented in Figures 6-8. Sequences were determined using standard techniques, which are described in the Exemplification.

10 V. Filter hybridization of Insert DNA

Arrays of lambda gt11 clones were created as described above and incubated at 42° for 5 hours. The plates were then overlayed with nitrocellulose filters and placed at 4°C for 1 hour. Probe DNA was labelled with ³²P by nick translation. Filter 15 hybridization was performed as described by Davis et al. in Advanced Bacterial Genetics: A Manual for Genetic Engineering, Cold Spring Harbor (1980), the teachings of which are incorporated herein by reference. Hybridization conditions were as follows: 50% 20 v/v formamide, 5x SSPE (1x SSPE is 0.18M NaCl, 10mM Na_{1.5}H_{1.5}PO₄, 1mM Na₂ EDTA, pH 7.0), 1x Denhardt's solution (0.02% w/v Ficoll, 0.02% w/v polyvinyl-pyrrolidone, 0.02% w/v bovine serum albumin), 0.3% NaDodSO₄ at 42°C for approximately 16 hours, followed by washing in 2x SSPE, 0.2% NaDodSO₄ at 45°C. 25

VI. Recombinant Antigens Recognized by Rabbit Serum

The response of a second animal to an antigen preparation of M. tuberculosis was assessed by

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examining the reactivity of rabbit anti-M. tuberculosis hyperimmune sera with recombinant antigens. Cloned lambda gt11 recombinants were arrayed on lawns of E. coli and probed with the rabbit hyperimmune serum. Anti-M. tuberculosis hyperimmune serum, produced by repeated immunization of rabbits with M. tuberculosis H37Rv culture filtrate, was provided by J. Bennedsen (Statens Serum Institut, Copenhagen, Denmark). These sera were used at 1:100 dilution.

These sera produced positive signals with lambda gt11 clones encoding each of the five M. tuberculosis epitopes which had been isolated with murine monoclonal antibodies (Figure 2). Particularly strong signals were observed with the 65kD and 71kD antigens (Figure 2). These results demonstrate that mice and rabbits can mount an antibody response to the same protein antigens of M. tuberculosis.

Clones for the five M. tuberculosis antigens were detected at similar frequencies in the lambda gt11 recombinant DNA library. Thus, the number and type of antigen-producing clones isolated with polyclonal serum antibodies should reflect the relative titer and diversity of the individual antibodies in this serum.

To determine whether any of the 5 M. tuberculosis antigens are relatively immunodominant in the rabbit humoral immune response to M. tuberculosis, the M. tuberculosis lambda gt11 recombinant DNA library was screened with the rabbit serum. Forty signal-producing clones were isolated, arrayed on

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lawns of E. coli Y1090 and probed with monoclonal antibodies directed against each of the 5 recombinant M. tuberculosis protein antigens. Remarkably, 17 of the 40 clones (43%) reacted strongly with at least one of the four anti-65kD monoclonal antibodies tested. An additional six clones (15%) reacted strongly with the anti-17kD monoclonal antibody (IT-11). This indicates that a large proportion of the anti-M. tuberculosis antibody present in the rabbit serum was directed against the 65kD antigen of M. tuberculosis and suggests that it is a dominant antigen for the rabbit humoral immune response. Seventeen of the clones did not react with any of the monoclonal antibodies tested, suggesting that the rabbit sera may identify M. tuberculosis proteins not recognized by the murine antibodies.

VII. Antigenic Relatedness of M. tuberculosis and M. leprae Proteins

There is evidence that M. tuberculosis and M. leprae share immunologically important antigens. To assess this further, an investigation of the exact nature of the immunological relatedness among recombinant protein antigens of M. tuberculosis and M. leprae was conducted.

For each of five M. tuberculosis and four M. leprae protein antigens, a single recombinant DNA clone containing most or all of the gene of interest was used to express antigen in the following manner. The recombinant phage clones were arrayed on a lawn

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of E. coli Y1090, which was then grown and induced for antigen expression.

Antigen immobilized on nitrocellulose filters was then probed with 26 individual anti-M. tuberculosis and M. leprae monoclonal antibodies. Figure 3 shows the array of DNA clones used and the results obtained with the anti-M. tuberculosis antibodies IT-1, IT-3, IT-11, IT-16, and IT-31, which recognized proteins of 14kD, 12kD, 71kD, 19kD and 65kD respectively. Table 1 details the full results of these cross-screening experiments, showing the reactivity of antigen expressed from individual recombinant DNA clones with each of the individual monoclonal antibodies. Clones were scored as positive only if the signal produced was clearly greater than the background signal produced by the non-recombinant lambda gt11 clone included in each array.

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TABLE 1
Reactivity of Monoclonal Antibodies with
Recombinant Protein Antigens

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Several conclusions can be drawn from the results shown in Table 1. Among the 11 monoclonal antibodies that recognize a 65kD antigen, 7 react with the 65kD protein from both mycobacteria (IT-31, 05 C1.1, IIH9 (identical to IT-33), IIC8, T2.3, Y1-2, SA2.D7C), one antibody reacts only with the M. tuberculosis 65kD protein (IT-13), and two antibodies react only with the M. leprae 65kD protein (IIIE9 and IIIC8). One antibody, ML30A, 10 cross-reacts with an antigen in E. coli and could not specifically identify antigen-producing clones. These results indicate that the 65kD protein antigens of M. tuberculosis and M. leprae are homologues and share a number of epitopes. In 15 addition to these shared epitopes, however, both 65kD antigens have epitopes that are specific for one species relative to the other.

No cross-reactivity was observed between other antigens of these two mycobacterial species. 20 Because monoclonal antibodies recognize a single epitope and because only one or a few antibodies were available for each antigen, it is not clear whether the 65kD proteins are the only homologous protein antigens of M. tuberculosis and M. leprae. 25 Among the antigens for which lambda gt11 clones have been isolated, the 18kD antigen of M. leprae and the 19kD antigen of M. tuberculosis are of similar size. To determine whether these two antigens are related, the homology of the DNA sequences that encode these 30 antigens was examined. At conditions of moderate stringency, no hybridization was observed between

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the insert DNA and Y3147 (an M. tuberculosis 19kD clone) and Y3179 (an M. leprae 18kD clone). This indicates no significant homology between the DNA sequences of the insert DNAs of these two clones.

05 This result suggests that the M. tuberculosis 19kD and the M. leprae 18kD proteins are unlikely to be homologous.

As a result of the work described, recombinant DNA clones encoding five major protein antigens of M. tuberculosis were isolated through the use of an extensive collection of well-characterized murine monoclonal antibodies. These five proteins were also found to be major antigens in the rabbit humoral immune response to M. tuberculosis. One of 15 these antigens, the 65kD protein, is shared with another other mycobacterial pathogen M. leprae.

Several lines of evidence indicate that the 65kD antigen is among the most immunodominant of the protein antigens of M. tuberculosis. Eleven of the 20 25 different M. tuberculosis and M. leprae monoclonal antibodies examined in this study recognized the 65kD recombinant antigen from one or both mycobacteria. In addition, almost half of the recombinant DNA clones isolated with rabbit poly- 25 clonal anti-M. tuberculosis sera express the 65kD antigen, reflecting the predominance of antibody to this antigen in these sera.

Considerable evidence indicates that the 65kD antigen plays an important role in the human response to tuberculosis. Antibodies directed against 30 this protein can be detected in the serum of

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patients with tuberculosis. The 65kD antigen is present in purified protein derivatives (PPD's) of M. tuberculosis, M. bovis, and other mycobacteria. Thole, J.E.R. et al., Infection Immunity, 50:800-806 05 (1985). Finally, helper T cell clones reactive with recombinant 65kD antigen have been isolated from patients with tuberculosis, indicating that this antigen is involved in the cell-mediated as well as the humoral immune response to tuberculosis.

10 Among the major antigens of the leprosy bacillus, the 65kD antigen appears to elicit antibody and T cell responses similar to those observed for the M. tuberculosis antigen. Both serum antibodies and T cells directed against the 65kD M.

15 leprae antigen have been observed in patients with leprosy. Britton, W.J. et al., Journal of Immunology, 135:4171-4177 (1985); Mustafa, A.S. et al., Nature, 319:63-66 (1986). In addition, T cell clones from leprosy patients have been found to

20 respond to recombinant 65kD protein of M. bovis, as well as to PPD's from both M. bovis BCG and M.

25 leprae. Emmrich, F. et al., Journal of Experimental Medicine, 163:1024-1029 (1986); Shankar, P. et al., Journal of Immunology, 136:4255-4263 (1986). It is interesting to note that in vaccine trials in Asia and Africa, BCG provided significant protection against leprosy, ranging from 20% to 80%. Fine, P., Tubercle, 65:137-153 (1984). An intriguing possibility is that the M. bovis BCG 65kD antigen is

30 involved in engendering the immune protection

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provided by this vaccine against M. leprae, as well as against M. tuberculosis.

In addition to the 65kD antigen, there is evidence that the 19kD and 71kD antigens of M. tuberculosis may be particularly important in the immune response to this bacillus. Helper T cell clones from tuberculosis patients have been isolated which respond to the recombinant 19kD protein. The 71kD antigen is recognized by the humoral immune system of both mice and rabbits, and antibody to this antigen has been shown to be a prominent component of hyperimmune anti-M. tuberculosis rabbit sera.

15 VIII. Isolation of DNA Clones for Genes Encoding Proteins of Additional Mycobacteria

Using the procedures described above for isolation of genes encoding M. tuberculosis proteins, genes encoding proteins of additional mycobacteria were isolated. DNA clones containing genes encoding the following proteins were isolated:

<u>Mycobacterium</u>	<u>Protein</u>	<u>Clone</u>
<u>M. bovis</u> BCG	71kD	PL1-101
	65kD	PL1-105
	19kD	PL1-501
	14kD	PL1-502
<u>M. smegmatis</u>	65kD	PL1-206
<u>M. avium</u>	65kD	PL1-401
<u>M. africanum</u>	65kD	PL1-301

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For purposes of comparison, genes encoding the following proteins were isolated for M. tuberculosis and M. leprae:

	<u>Mycobacterium</u>	<u>Protein</u>	<u>Clone</u>
05	<u>M. tuberculosis</u>	71kD	Y3272
		65kD	Y3150
		19kD	Y3147
		14kD	Y3248
	<u>M. leprae</u>	65kD	

10 The following strains were used for this purpose:

	<u>Species</u>	<u>Isolate</u>
	<u>M. leprae</u>	Armadillo isolate (WHO)
15	<u>M. tuberculosis</u>	Erdmann strain
	<u>M. africanum</u>	African clinical isolate
	<u>M. bovis</u> BCG	Danish vaccine strain
	<u>M. smegmatis</u>	MC ² -6
	<u>M. avium</u>	AIDS patient isolate

20 DNA from recombinant lambda gt11 clones was isolated, as described above, and mapped with restriction endonucleases, using standard techniques. Davis, R.W. et al., Advanced Bacterial Genetics: A Manual for Genetic Engineering, Cold Spring Harbor (1980).

25 Figure 4 presents a comparison of the restriction maps for four genes of M. tuberculosis with the restriction maps for four genes of M. bovis BCG which encode proteins of the same molecular weight. As is evident from the figure, in each case, the

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restriction sites on the two genes (e.g., those on the M. tuberculosis gene and those on the M. bovis gene which encodes a protein of the same molecular weight) are essentially identical. This indicates
05 that the sequence of the genes of the two mycobacteria (at least those encoding these four proteins) are very similar and, therefore, the proteins they encode are also very similar.

Figure 5 presents a comparison of the restriction map for the gene encoding the 65kD protein for the six mycobacteria. As is evident, the restriction maps for the genes encoding the 65kD protein of M. tuberculosis, M. africanum, M. bovis BCG, M. smegmatis and M. avium are essentially identical.
10 The fact that there is no detectable difference among these mycobacteria at the level of the restriction map is an indication that, at least at this level, the encoded proteins are the same.
15

As is also evident, the map of the M. leprae 20 65kD gene has several identical restriction sites in common with those of the other mycobacteria; it also has two sites not found in the other genes and lacks three sites present in the others. This indicates that, at the level of the restriction map, there are 25 similarities in the DNA (and the encoded protein). In addition, however, there are differences apparent at this level.

IX. Diagnostic, Therapeutic and Preventive Applications

30 The isolation of genes encoding major protein antigens of M. tuberculosis makes it possible to

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address problems which presently exist in diagnosing
treating and preventing tuberculosis. Isolation of
genes encoding proteins of other mycobacteria, such
as M. bovis BCG, M. africanum, M. smegmatis and M.
05 avium makes it possible to address similar problems
in diseases which they cause.

The nucleotide sequence of three of the five
genes has been determined. The sequence of the
remaining genes can be determined using well-known
10 methods, such as that of Sanger et al. Sanger, F.
et.al., Proceedings of the National Academy of
Sciences, USA, 74:5463-5467 (1977). The amino acid
sequence of each of the immunodominant proteins has
been deduced from the nucleotide sequence of the
15 three genes and can be done for the others.

Identification and characterization of the
genes for major tuberculosis protein antigens and of
the proteins themselves make it possible to develop
improved reagents for diagnosis and immuno-
20 prophylaxis of tuberculosis. Proteins antigens
encoded by an entire gene, or amino acid sequences
(e.g., peptides, protein fragments) which make up
the antigenic determinant of a M. tuberculosis
25 antigen (i.e., M. tuberculosis-specific antigenic
determinants) may be used in serodiagnostic tests
and skin tests. Such antigens would be highly
specific to the tuberculosis bacillus and the tests
in which they are used would also be highly
specific. Highly specific serological tests would
30 be of great value in screening populations for

-26-

individuals producing antibodies to M. tuberculosis-specific antigenic determinants; in monitoring the development of active disease in individuals and in assessing the efficacy of treatment. As a result,
05 early diagnosis of tuberculosis will be feasible, thus making it possible to institute treatment at an early stage of the disease and, in turn, to reduce the likelihood it will be transmitted.

As a result of the work described, it is also
10 possible to determine which segment(s) of the M. tuberculosis antigen is recognized by M. tuberculosis-specific T cells. A mixture of peptides recognized by helper T cells can serve as a specific skin test antigen useful in assessing
15 immunological status (delayed hypersensitivity) of infected individuals and those with whom they come in contact. This specific skin test antigen would be useful in evaluating rapidly the immunological efficacy of anti-tuberculosis vaccines.

20 It is reasonable to expect that the products encoded by M. tuberculosis genes, particularly those shown to be recognized by helper T cells, are themselves immunogenic and thus useful components of vaccines against tuberculosis. These products
25 include proteins and portions of such proteins (e.g., polypeptides and peptides). For example, one approach to vaccine development is the introduction of genes encoding products (e.g., polypeptides) which provide immunological protection into viruses
30 such as vaccinia virus, or bacteria, such as cultivatable mycobacteria, thus producing a vaccine

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capable of engendering long-lasting and very specific immunity. The genes encoding five immunodominant protein antigens of the tuberculosis bacillus, described herein, are useful for that 05 purpose; genes encoding the 65kD, 19kD and 71kD antigens, or a portion thereof, are particularly valuable in vaccine construction.

Because of the similarities in the DNA encoding similarly-sized proteins and, thus, of the encoded 10 proteins themselves, it is possible that, for example, a vaccine effective against two or more of the mycobacteria can be produced.

EXEMPLIFICATION

Isolation and Analysis of Recombinants Expressing the 65kD M. tuberculosis Antigen

The recombinant DNA library of M. tuberculosis genomic DNA fragments in the lambda gt11 vector was constructed as described above. Recombinant phage lambda RY3143 and lambda RY3146 were used. Young, 20 R.A. et al., Proceedings of the National Academy of Sciences, USA, 82:2583-2587 (1985). Subclones of the mycobacterial DNA inserts in these recombinant phage were constructed in pUC19 or M13mp9 vectors using standard recombinant DNA techniques. Messing, J. and J. Viera, Gene, 19:269-276 (1982). Maniatis, T. et al., Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory, Cold Spring Harbor, 25 NY (1982).

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Monoclonal antibodies specific for the 65kD antigen were obtained from the Immunology of Tuberculosis Scientific Working Group under a grant from the WHO/World Bank/UNDP Special Program for 05 Vaccine Development. These antibodies included IT-13 (WTB-78), IT-31 (SA2D5H4), and IT-33 (MLIIH9). Coates, A.R.M. *et al.*, Lancet, 2:167-169 (1981). Gillis, T.P. and T.M. Buchanon, Immunology, 37:172-178 (1982). Anti-B-galactosidase antibodies 10 were purchased from CooperBiomedical. Polyclonal rabbit antisera directed against a sonicate of M. tuberculosis strain H37Rv were elicited as described by Minden and co-workers. Minden, P. *et al.*, Infect. Immun., 46:519-525 (1984). Results are 15 shown in Table 2.

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TABLE 2: PATTERNS OF ANTIBODY REACTIVITIES^a

	<u>Number of Clones</u>	<u>Reactivity with Antibodies</u>		
		<u>IT-13</u>	<u>IT-31</u>	<u>IT-33</u>
05	27	+	+	+
	1	+	+	+
	2	+	-	+
	3	-	+	+
	1	+	-	-
	2	-	+	-
10	2	-	-	+

^a: Recombinant clones expressing antigens reactive with the 65kD antigen specific monoclonal antibodies IT-13, IT-31, and IT-33 were isolated as described above. For the initial screen, a pool of the three antibodies was used; it contained a 1:1000 dilution of each antibody to screen a total of about 8×10^5 recombinant phage from the lambda gt11-M. tuberculosis library. To determine which monoclonal antibody reacted with which of the 38 plaque-purified recombinants, about 100 pfu of each recombinant phage were inoculated in small spots on a lawn of Y1090. The phage were allowed to grow and induced to synthesize the foreign proteins as described previously. The filters were then reacted with a 1:1000 dilution of one of the monoclonal hybridoma antibodies as described above.

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The lambda gt11-M. tuberculosis library was screened with the monoclonal antibodies specific for the 65kD antigen and clones reactive with them were isolated essentially as described by Young et al.

05 Young, R.A. et al., Proceedings of the National Academy of Sciences, USA, 82:2583-2587 (1985). Briefly, for each 150mm LB plate, 0.6ml of a fresh overnight culture of Y1090 was infected with 1-2 $\times 10^5$ plaque forming units of the library. After
10 3.5-4 hours of growth at 42°C, the plaques were overlaid with a dry nitrocellulose filter which had been saturated with 10mM isopropyl-B-D-thiogalactopyranoside (IPTG). The plates were incubated an additional 3.5-4 hours at 37°C and then removed to
15 room temperature and the position of the filters marked. The filters were washed briefly in TBST (50 mM Tris-HCl, pH 8, 150mM NaCl, 0.05% Tween 20) and then incubated in TBST + 20% fetal calf serum. After 30 minutes at room temperature, the filters
20 were transferred to TBST plus antibody. For the initial screen, the antibody mix contained a 1:1000 dilution of IT-13, IT-31, and IT-33. The filters were incubated with the antibody solution overnight at 4°C with gentle agitation, washed in TBST and reacted with biotinylated goat anti-mouse immuno-globulin, the Vectastain ABC reagent, and developer as described by the manufacturer (Vector Laboratories). After the color had developed the filters were washed with several changes of water
25 and air dried. Phage corresponding to positive signals were twice plaque purified. To determine
30

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which monoclonal antibodies reacted with which of
the recombinant phage, about 100 pfu of a purified
phage stock were inoculated in a small spot on a
lawn of Y1090 bacteria on an LB plate. The phage
05 were allowed to grow and induced to synthesize the
foreign proteins as described above. The filters
were then reacted with a 1:1000 dilution of one of
the monoclonal antibodies. The subsequent steps
were the same as for the initial screen.

10 Western blot assays were carried out as
follows: Cells containing phage or plasmids in
which the expression of the foreign sequences was
under the control of the E. coli lac gene regulatory
sequences were induced to synthesize the foreign
15 proteins by incubating the cells in the presence of
2.5mM IPTG for 2 hours. Crude lysates of cells
expressing lambda gt11 recombinants were made as
described in Huynh et al. Huynh, T.V. et al., In:
DNA Cloning Techniques: A Practical Approach, (D.
20 Glover, ed.) IRL Press, Oxford, Vol. 1, pp. 49-78
(1985). Crude lysates of cells expressing plasmid
encoded proteins were made by harvesting cells from
overnight cultures and resuspending the cells in 10
mM Tris pH7.5/10 mM EDTA containing 100 ug
25 lysozyme/ml. After 10 minutes at room temperature,
SDS was added to a final concentration of 0.5%. A
protease inhibitor (Trasylol, Boehringer Mannheim)
was added to all crude lysates at a final concen-
tration of 0.3%. The crude protein preparations
30 were electrophoresed on 10% polyacrylamide-SDS
Laemmli gels and the separate proteins electrophor-

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etically transferred to nitrocellulose. Laemmli,
U.K., Nature, 227:680-685 (1970). Towbin, H. et
al., Proceedings of the National Academy of
Sciences, USA, 76:4350-4354 (1979). The immobilized
05 proteins were reacted with a 1:1000 dilution of
monoclonal antibody IT-13 in TBST overnight at 4°C.
The nitrocellulose filters were then washed, reacted
with peroxidase-conjugated goat anti-mouse immuno-
globulin, and developed as described by Niman and
10 co-workers. Niman, H.L. et al., Proceedings of the
National Academy of Sciences, USA, 80:4949-4953
(1983).

The sequences of 5'-end-labeled restriction
fragments of the mycobacterial DNA were determined
15 by a modification of the partial chemical
degradation technique of Maxam and Gilbert. Brow,
M.A.D. et al., Mol. Biol. Evol., 2:1-12 (1985).
Maxam, A.M. and W. Gilbert, Proceedings of the
National Academy of Sciences, USA, 74:560-564
20 (1976). For the M13/dideoxy sequencing studies,
Sau3AI fragments from the mycobacterial DNA inserts
were subcloned into the BamHI site of M13mp9. Phage
DNA was isolated from the M13 recombinants and
subjected to the dideoxy chain termination
25 sequencing reactions. Biggin, M.D. et al.,
Proceedings of the National Academy of Sciences,
USA, 80:3963-3965 (1983). Sanger, F. et al.,
Journal of Molecular Biology, 143:161-178 (1980).
The products of the sequencing reactions were
30 electrophoresed on 6% acrylamide/7M urea/0.5-2.5 x
TBE gradient sequencing gels. The gels were dried

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under vacuum and exposed to Kodak XRP-1 film. The nucleotide sequences were determined independently for both strands of the mycobacterial DNA.

Computer-aided analyses of the nucleic acid sequences and deduced protein sequences were performed using the Databases and programs provided by the Nucleic Acid and Protein Identification Resources of the National Institutes of Health as well as the programs of Chou and Fasman and Hopp and Woods. Chou, P.Y. and G.D. Fasman, Adv. Enzym., 47:45-148 (1978). Hopp, T.P. and K.P. Woods, Proceedings of the National Academy of Sciences, USA, 78:3824-3828 (1981). The nucleotide sequence of the region containing the M. tuberculosis 65kD gene and the deduced amino acid sequence of the two long open reading frames are represented in Figure 8.

B-galactosidase assays were also carried out. Cells were grown in LB broth or LB broth plus 2.5mM IPTG to an OD₆₀₀ of about 0.3. Crude lysates were made and b-galactosidase activity assayed as described by Miller. Miller, J.H., Experiments in Molecular Genetics, Cold Spring Harbor Laboratory, Cold Spring Harbor, NY (1972).

25 Equivalents

Those skilled in the art will recognize or be able to ascertain, using no more than routine experimentation, many equivalents to the specific materials and components described herein. Such equivalents are intended to be encompassed in the scope of the following claims.

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CLAIMS

1. Isolated DNA encoding an immunogenic protein antigen of Mycobacterium tuberculosis.
2. DNA of Claim 1 selected from the group consisting of DNA encoding Mycobacterium tuberculosis protein antigens of molecular weight 71kD, 65kD, 19kD, 14kD and 12kD.
3. Isolated DNA encoding an immunodominant protein antigen of Mycobacterium tuberculosis, the protein antigen having a molecular weight of approximately 65kD and recognized by a monoclonal antibody selected from the group consisting of: IT-31; C1.1; IIH9; IIC8; T2.3; Y1-2; SA2.D7C and IT-13.
4. Isolated DNA encoding an immunodominant protein antigen of Mycobacterium tuberculosis, the protein antigen having a molecular weight of approximately 19kD and recognized by a monoclonal antibody selected from the group consisting of: IT-10; IT-12; IT-16; and IT-19.
5. Isolated DNA encoding an immunodominant protein antigen of Mycobacterium tuberculosis, the protein antigen having a molecular weight of approximately 71kD and recognized by the monoclonal antibody IT-11.

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6. Isolated DNA encoding an antigenic determinant of Mycobacterium tuberculosis protein.
7. DNA of Claim 6 which encodes an antigenic determinant selected from the group consisting of antigenic determinants of Mycobacterium tuberculosis proteins of molecular weight 71kD, 65kD, 19kD, 14kD and 12kD.
05
8. Isolated DNA encoding an amino acid sequence of an antigenic determinant of Mycobacterium tuberculosis protein, said protein having a molecular weight of approximately 65kD.
10
9. Isolated Mycobacterium tuberculosis DNA encoding an immunodominant protein antigen having a molecular weight of approximately 65kD, said DNA selected from the group consisting of:
15
 - a. the DNA insert of clone Y3141;
 - b. the DNA insert of clone Y3143;
 - c. the DNA insert of clone Y3150;
 - d. the DNA insert of clone Y3253; and
20
 - e. the DNA insert of clone Y3262.
10. A protein antigen encoded by DNA of Claim 9.
11. A protein antigen of Claim 10, wherein the protein antigen is recognized by a monoclonal antibody selected from the group consisting of
25 IT-31; C1.1; IIH9; IIC8; T2.3; Y1-2; SA2.D7C and IT-13.

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12. Isolated DNA having a nucleotide sequence selected from the group consisting of: a) the nucleotide sequence represented in Figure 6, or a portion thereof; b) the nucleotide sequence represented in Figure 7, or a portion thereof; and c) the nucleotide sequence represented in Figure 8, or a portion thereof.
13. A protein or a peptide selected from the group consisting of: a) proteins or peptides encoded by the nucleotide sequence represented in Figure 6, or a portion thereof; b) proteins or peptides encoded by the nucleotide sequence represented in Figure 7, or a portion thereof; and c) proteins or peptides encoded by the nucleotide sequence represented in Figure 8, or a portion thereof.
14. A peptide having the amino acid sequence of an antigenic determinant of Mycobacterium tuberculosis protein, said antigenic determinant being unique to Mycobacterium tuberculosis protein.
15. A peptide encoded by isolated Mycobacterium tuberculosis DNA, said peptide recognized by helper T cells.
16. A peptide encoded by the Mycobacterium tuberculosis DNA insert of clone Y3150 or a portion of said DNA insert.

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17. Isolated DNA encoding a protein of Mycobacterium africanum, the protein having a molecular weight of 65kD.
18. Isolated DNA encoding a protein of Mycobacterium avium, the protein having a molecular weight of 65kD.
05
19. A vaccine comprising DNA encoding Mycobacterium tuberculosis protein in a recombinant vaccine vector capable of expressing said DNA.
10. 20. A vaccine of Claim 19 in which the recombinant vaccine vector is vaccinia virus or cultivatable mycobacteria.
21. A vaccine of Claim 20 in which the DNA encodes the 65kD Mycobacterium tuberculosis protein recognized by the monoclonal antibody IT-13, or
15 a portion of said protein.
22. A vaccine comprising DNA encoding an antigenic determinant unique to Mycobacterium tuberculosis cultivatable mycobacteria capable of expressing said DNA.
20
23. A method of detecting antibody against Mycobacterium tuberculosis in a biological fluid, comprising the steps of:
25 a) incubating an immunoadsorbent comprising a solid phase to which is attached

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- 05 immunodeterminant Mycobacterium tuberculosis protein with a sample of the biological fluid to be tested, under conditions which allow the anti-Mycobacterium tuberculosis antibody in the sample to bind to the immunoadsorbent;
- 10 b) separating the immunoadsorbent from the sample; and
- 10 c) determining if antibody is bound to the immunoadsorbent, as an indication of anti-Mycobacterium tuberculosis in the sample.
24. A method of Claim 23 in which the Mycobacterium tuberculosis protein attached to the solid phase has a molecular weight of approximately 65kD.
- 15 25. A method of detecting antibody against Mycobacterium tuberculosis in a biological fluid, comprising the steps of:
- 20 a) incubating an immunoadsorbent comprising a solid phase to which is attached a peptide having the amino acid sequence of an antigenic determinant of Mycobacterium tuberculosis protein with a sample of the biological fluid to be tested, under conditions which allow antibody against Mycobacterium tuberculosis to bind to the immunoadsorbent;
- 25 b) separating the immunoadsorbent; and
- 25 c) determining if antibody is bound to the immunoadsorbent, as an indication of the

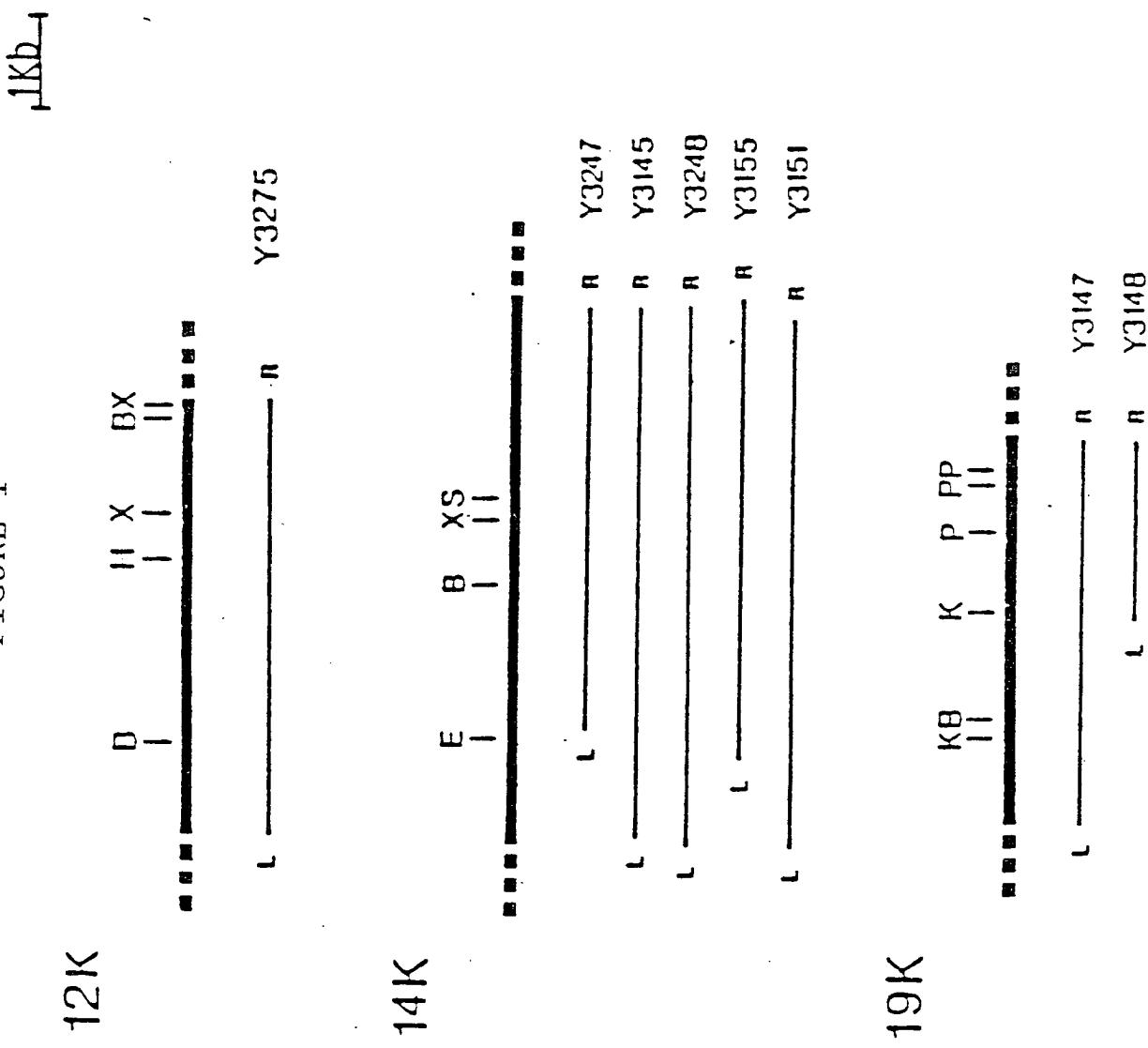
-39-

presence of the antibody against Mycobacterium tuberculosis in the sample.

- 05 26. A method of Claim 25 in which the peptide has the amino acid sequence of an antigenic determinant which is unique to Mycobacterium tuberculosis protein.
- 10 27. A kit useful in detecting antibody against Mycobacterium tuberculosis in a biological fluid, comprising a collection of reagents for immunoassay of said antibody, said collection of reagents a solid phase to which is attached immunodeterminant Mycobacterium tuberculosis protein or a peptide having the amino acid sequence of an antigenic determinant of Mycobacterium tuberculosis.
- 15

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FIGURE 1

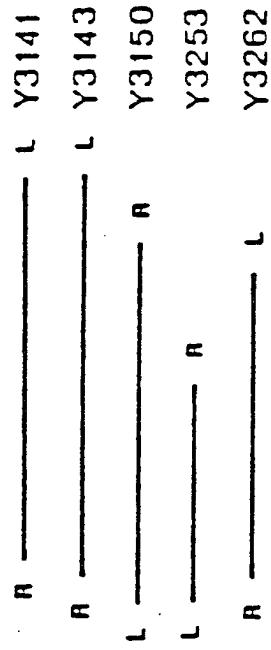
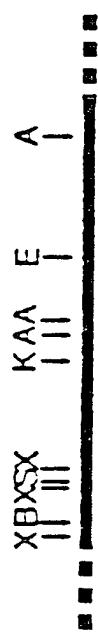


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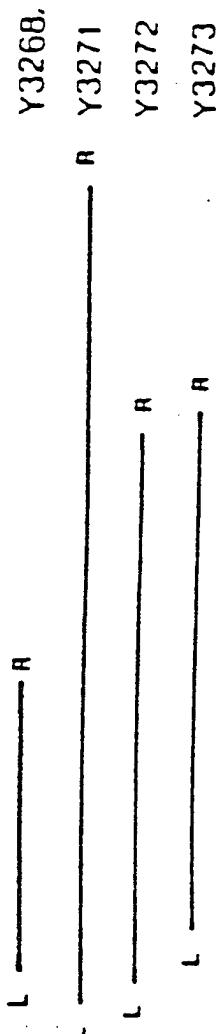
FIGURE 1 (Cont'd)

1Kb

65K

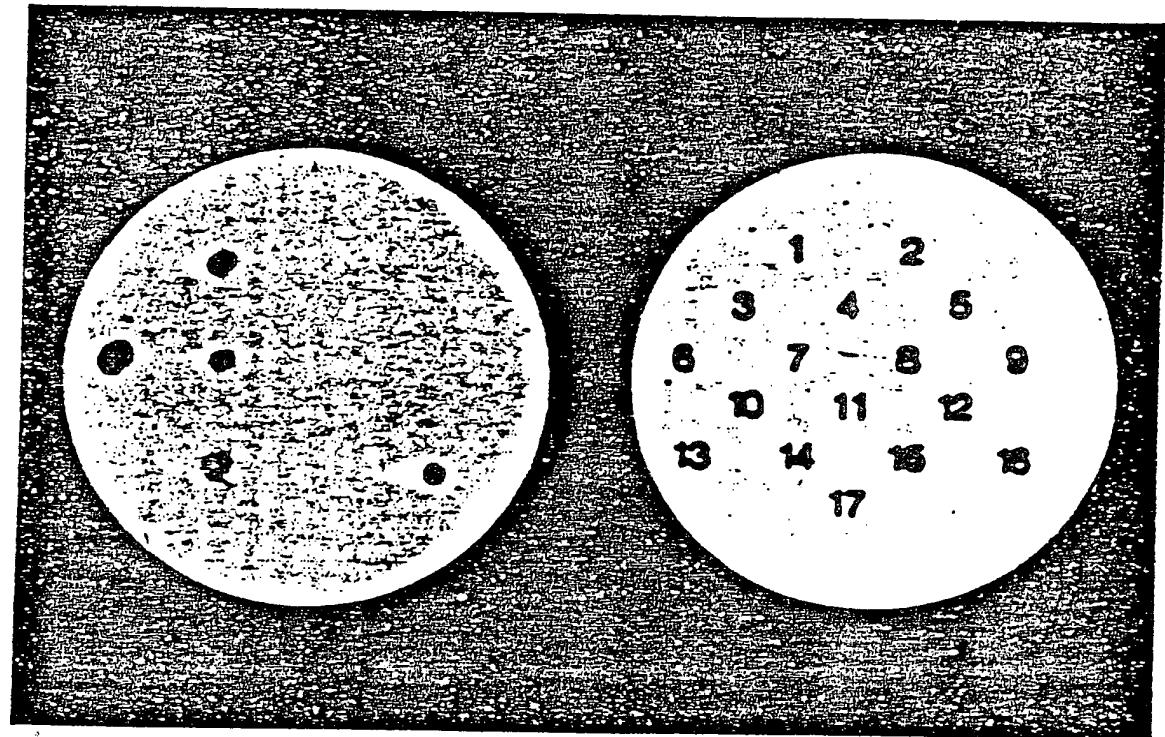


71K



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FIG.2



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FIG.3

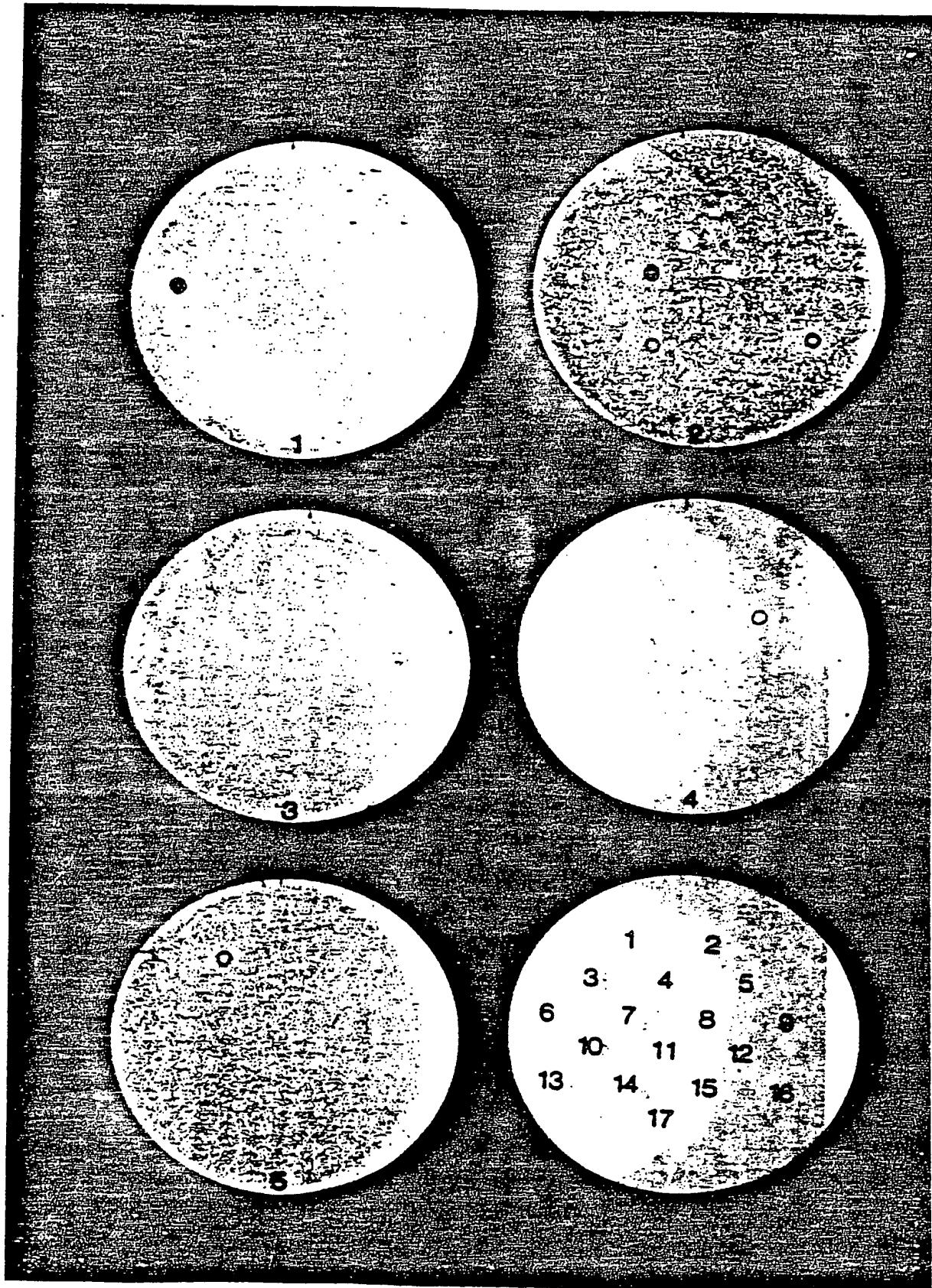
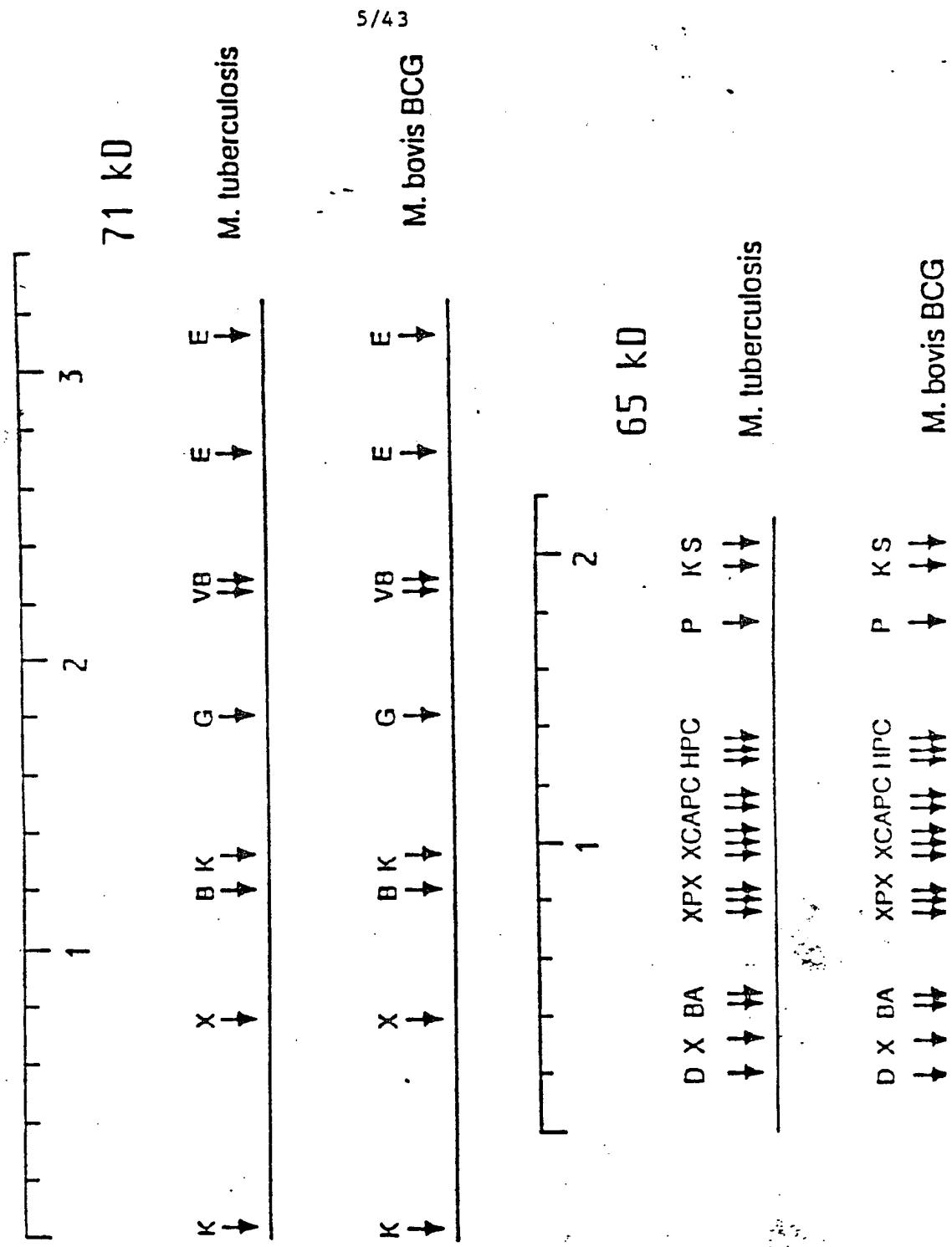


FIGURE 4

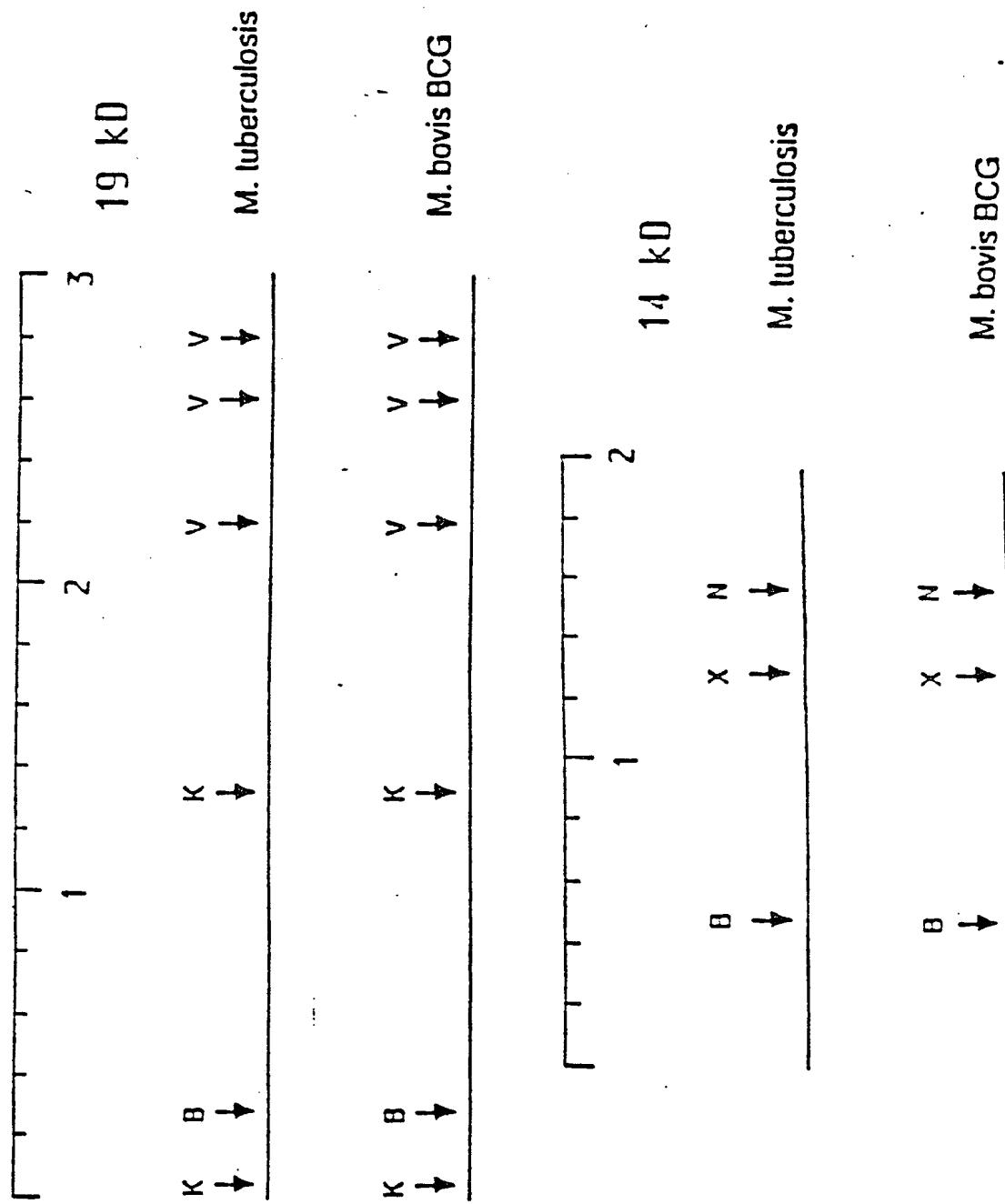


D X BA X PX X CAPC HPC P KS
 ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓
 ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓

M. bovis BCG

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FIGURE 4 (Cont'd)



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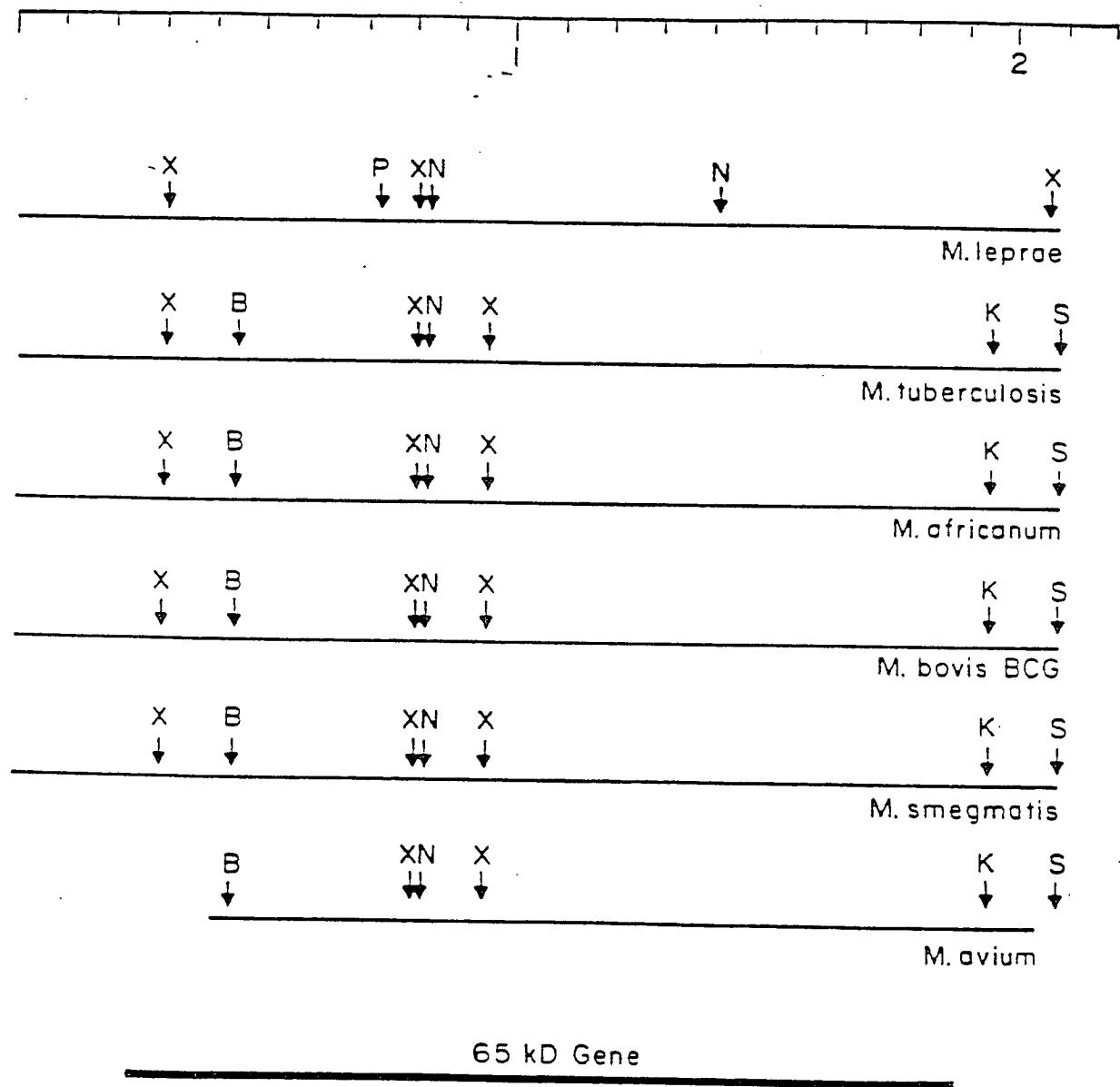


FIG. 5

FIGURE 6

F R V N R L G E I A R P G A R I A H Q G
 S V S I A S V R * P D Q A R G S R T K A
 P C Q S P R * D S P T R R A D R A P R R
 TTCCGTGTCATGGCTCGGTGAGATAAGCCGACCCAGGGATCGGCCACCAAGGC
 10 20 30 40 50 60
 AAGGCACAGTTAGGGAGCCACTCTATGGCTGGTCCGGCCTAGGGCGTGGTCCG
 E T D I A E T L Y G S W A R P D R V L A
 G H * D G R H S L G V L R A S R A G; L R
 R T L R R P S I A R G P A R I A C W P A
 A A Q G Q V V Q V M R G V F G H A Q R
 P R R V S R L C R L C V A F S A M H N A
 R A G S A G C A G Y A W R F R P C T T R
 GCCGGCAGGGTCAGCAGGGTTATGCCTGGCGTTTCGGCCATGCACAAACGC
 70 80 90 100 110 120
 CGGGCGGTCCCAGTCGTCCAACACGTCCAAATACGCACCGCAAAGCCGGTACGTGGCG
 G R L T L N H L N H T A N E A M C L A
 R A P D A P Q A P * A H R K R G H V V R
 A C P * C T T C T I R P T K P W A C R A

FIGURE 6 (CONT'D)

A E A R E I E V H L R R G L G A R G H L
 R K R G K * K C I S A G A S V P G A I W
 G S A G N R S A S P Q G P R C P G P S G
 GCGGAAGCGGGAAATAGAAGTGCATCTCCGAGGGCCATCTG
 130 140 150 160 170 180
 CGCCTTCGGCCCTTATCTCACGTTAGAGGGGTCCCCGGAGCCACGGGGCCAGGTAGAC
 R F R P F Y F H M E A P A E T G P A M Q
 P L A P F L L A D G C P G R H G P G D P 9/43
 S A R S I S T C R R L P R P A R P W R S
 E L D L H P V D G V G L P G L G D V D R
 N S I S T P S M V W V S P V S V M S T V
 T R S P P R R W C G S P R S R * C R P S
 GAACTCGATCTCCACCCCGTCGATGGTGTGGGTCTCGGTGATGTCCGACCGT
 190 200 210 220 230 240
 CTTGAGCTAGGGTGGGCAGCTACCCACAGGGCCAGAGCCACTACAGCTGGCA
 F E I E V G D I T H T E G T E T I D V T
 V R D G G R R H H P D G R D R H H R G D
 S S R W G T S P T P R G P R P S T S R R

FIGURE 6 (CONT'D)

R H D E R N L T G R Q C L P E A A A D V
 G T T S E T S P V D S V C P R P Q P T C
 A R R A K P H R S T V S A R G R S R R A
 CGGCACGGAGCCAAACCTCACCGGTGACAGTGTCTGCCAGGGCAGCCGACTG
 250 260 270 280 290 300
 GCCGTGCTCGCTTGGAGTGGCCAGCTGTCAACAGACGGGCTCCGGCTGGCAC
 P V V L S V E G T S L T Q G L G C G V H
 A R R A F G * R D V T D A R P R L R; R A
 C S S R F R V P R C H R G S A A S T G^{10/43}
 P P E T A R Q H G A V H V A R T A H H R
 P R R P R A N T V P Y M * P A R R I I A
 P G D R A P T R C R T C S P H G A S S P
 CCCCGGAGACCCGCCAACACGGTGGTACATGTAGCCCGACGGCATCATCGC
 310 320 330 340 350 360
 GGGGCCTCTGGCGGGTTGTGCACTGGGCATGTACATGGGGCATGGCG
 G R L G R A L V T G Y M Y G A R R M M A,
 G P S R A G V R H R V H L G C P A D D G
 G S V A R W C P A T C T A R V A C * R R

FIGURE 6 (CONT'D)

R A G V D V F L H G V R G E P L R R Q H
 E P A * M F S C T A C A V N P S G A S T
 S R R C F P A R R A R * T P P A P A P
 CGAGCCGGCGTAGATGTTCTGCACGGCGTGGTGAACCCCTCCGGGCCAGCAC
 370 380 390 400 410 420 430 440 450 460 470 480
 GCTCGGCCATCTACAAAGGACGTGCCAACGGCCACTTGGGAGGGCCGGCTG
 S G A Y I N E Q V A H A T F G E P A L V
 L R R L H K G A R R A R H V G G A G A G
 A P T S T K R C P T R P S G R R R W C R
 R H L S R V H V G L G G D A E H P T E M
 A T F P A S T S A W V V T P S T P P K *
 P P F P R P R R P G W * R R A P H R N D
 CGCCACCTTCCGGTCCACGTCTGGTGGTGAACGGCAGCACCCACCGAAATG
 430 440 450 460 470 480
 GCGGTGGAAAGGGCCAGGTGCAGCGGACCCACTGGCTCGTGGGGCTTAC
 A V K G A D V D A Q T T V G L V G G F H
 G G K G R G R R G P H R R A G W R F S
 W R E R T W T P R P P S A S C G V S I I

FIGURE 6 (CONT'D)

I D M A V G V D D R D H G A V G S A V G
 S T W L W V * M T A T T G R S A P R W A
 R H G C G C R * P R P R G G R L R G G R
 ATCGACATGGCTGTGGTAGATGACCGGGACCAACGGGGTGGCTCCGGGTGGC
 490 500 510 520 530 540 541^{12/43}
 TAGCTGTACGGACACCCACATCTACTGGGCTGGTGGCCAGGCCAGGGCGCCACCCG
 D V H S H T Y I V A V V P R D A G R H A
 R C P Q P H L H G R G R P P R S R P P R
 S M A T P T S S R S W P A T P E A T P A
 A I Q V Q R G G H L G G H Q R V D D D
 R Y K S S A A A T S V D T N G S M T I
 D T S P A R R P R W T P T G R * R S
 GCGATAAGTCCAGCGGGGGGCCACCTCGGTGGACACCAACGGGTGGATGACGAT
 550 560 570 580 590 600
 CGCTATGTTCAAGGTGGTCGGCGCCGGGTGGAGCCACCTGTGGTGGCTACTGCTA
 R Y L D L A A A V E T S V L P D I V I
 S V L G A R R R G G R H V G V P R H R D
 I C T W R P P W R P P C W R T S S S *

FIGURE 6 (CONT'D)

Q P S V T L N E A D I G D I E S A D L I
 S P V S P S T K L I L E I S N P R T * *
 A Q C H P Q R S * Y W R Y R I R G P D R
 CAGCCAGTGTCAAC CCTCAACGAAGATATTGGAGATA TCCGGGACCTGATA
 610 620 630 640 650 660
 GTCGGGTACAGTGGAGTTGCCTGACTATAACCTCTATAAGCTTAGGCCCTGGACTAT
 L G T D G E V F S I N S I D F G R V Q Y
 A W H * G * R L Q Y Q L Y R I R P G S L
 G L T V R L S A S I P S I S D A S R I S
 13/43
 D A R H H L V E A L F R G Q L G L P P Q
 M P G T T W * R P C F A V S W D C R H R
 C P A P P G R G P V S R S A G I A A T G
 GATGCCGGCACACCTGGTAGAGGCCCTGTTGGCTGAGCTGGGATTCGGCCACAG
 670 680 690 700 710 720
 CTACGGCCGTGGACCATCTCCGGACAAAGCGCCAGTCGACCCTAACGGGGTGT
 I G P V V Q Y L G Q K A T L Q S Q R W L
 II G A G G P L P G T E R D A P I A A V P

FIGURE 6 (CONT'D)

A R C W R T S A R N R P * S P N G G C A
 A G M H R C R R G T V E K R V R V V P
 L G C T D V G A A P S R N E Y A S L S H
 W D A P M S A R H R R E T S T R R C P T
 GCTGGATGCACCGATGTCGGCGGCCACCGTCGAGAACGAGTACGGTGTCCCCA
 730 740 750 760 770 780
 CGACCCTACGGCTACAGCCGGCCGGCTCTTGCTCATGGCAGCAACAGGGT
 S P H V S T P A A G D L F S Y A D N' D W
 Q S A G I D A R C R S V L V R R Q G V
 P I C R H R R P V T S F R T R T G C⁷⁴³
 H H A T I G S L D H T R G Q R G N E S A
 T T R P S A A L I T H G D S A A M N P R
 P R D H R Q P * S H T G T A R Q * I R D
 CACCAACGGACCATCGGCAGCCATTGATCACACACGGGACAGGGCAATGAATCCGGC
 790 800 810 820 830 840
 GTGGTGGCTGGTAGGCCGGAACTAGTGTGGCCCTGTGGCGCGTTACTTAGGGCGC
 V V R G D A K I V C P S L A A I F G R
 G R S W R C G Q D C V P V A R C H I R S
 W A V M P L R S * V R P C R P L S D A I

FIGURE 6 (CONT'D)

I G V V E I R C V M Q R * R V F T V C R
 S A S S K S V V S C N G N E C S P C A A
 R R R R N P L C H A T V T S V H R V P P
 ATCGGGCGTCGAAATCCGGTTGTCAACGGTAACGAGTGTGCCCCG 15/43
 850 860 870 880 890 900
 TAGCCGGAGCAGCTTACGGCAACACAGTAGCTGCATTGCTCACAAAGTGGCACACGGCG
 D A D D F D T T D H L P L S H E G H A A
 R R R R F G N H * A V T V L T * R T G G
 P T T S I R Q T M C R Y R T N V T H R R
 L D D G S G R F V F H R H Y I A T T T V
 W M T A V G G L C S I G T T L P L L R C
 G * R Q W E V C V P S A L H C H Y Y G A
 CTGGATGACGGCAGTGGAGGTTGTGTTCCATGCCACTACATTGCCACTACGGTG
 910 920 930 940 950 960
 GACCTACTGCCGTACCCCTCCAAACAAAGGTAGCCGTGATGATGCCAC
 Q I V A T P P K H E M P V V N G S S R H
 P H R C H S T Q T G D A S C Q W * * P A
 S S P L P L N T N W R C * M A V V V T C

FIGURE 6 (CONT'D)

H A G R C R W R T T L P T R K R E F S A
 T P V D A V G E P R Y R P E R E N F P P
 R R * M P L A N H A T D Q K E R I F R R
 CACGCCGGTAGATGCCCGTGGCAACCACGCTACCGACCAGAATTTCGGCC
 970 980 990 1000 1010 1020
 16/43
 GTGCCGGCATCTACGGCAACCGCTTGGTGGATGGCTCTCTCTAAAGGCCG
 V G T S A T P S G R * R G S L S F ,K G G
 R R Y I G N A F W A V S W F S L I K R R
 A P L H R Q R V V S G V L F L S N E A A
 A P R P R A L L T R I L P K R S S M P M
 H L D L G P C * R A Y C R S G P Q C R W
 T * T S G P A N A H T A E A V L N A D G
 GCACCTAGACTGGGCCCTGCTAACGGCATACTGCCAGCGTCCATGCCGATG
 1030 1040 1050 1060 1070 1080
 CGTGGATCTGGAGCCGGACGATTGGCGTATGACGGCTTGCAGGAGTTACGGCTAC
 C R S R P G Q * R A Y Q R L P G * H R H

FIGURE 6 (CONT'D)

V * V E P G A L A C V A S A T R L A S P
G L G R A R S V R M S G F R D E I G I S

D R Y D R Q R S T G * S V D * R S R * P
T A T T G K G A Q G E A W T D G R G S R
P L R Q A K E H R \int_V K R G L T V A V A G
GACCGCTACGGCACAGGAAAGGAGCACAGGGTGAAGCGTGGACTGACGGTAGGCC
1090 1100 1110 1120 1130 ; 1140
CTGGCGATGCTGTCCGTTCCCTCGCACCTGACTGCCAGGCCATGGC
V A V V P L P A C P S A H V S P R P L R
G S R C A F S C L T F R P S V T A T A P
R * S L C L L V P H L T S Q R D R Y G S

E P P F W S Q V F P D V Q A T S R L Q E
S R H S G R R S F R M F K Q Q V D Y R K
A A I L V A G L S G C S S N K S T T G S
GAGCCGCATTCTGGTCGGTCTTCGGATGTTCAAGCAACAAGTCGACTACAGGAA
1150 1160 1170 1180 1190 1200
CTCGGGGTAAAGACCAGCGTCCAGGAAGGCCTACAAAGTTCTGTGTCAGCTGATGTCCTT
L R W E P R L D K R I N L C C T S * L F
A A M R T A P R E P H E L L D V V P L
R N O N R T K G S T * A V L R S C S A

FIGURE 6 (CONT'D)

A V R P R P R Q A R R Q A P A P P P G R
 R * D H D R G R H D G K P R R R L R A E
 G E T T A A G T T A S P G A A S G P K
 GCGGTGAGACCACGACCGGGCACGGCAAGCCGCCTCCGGCCGA
 1210 1220 1230 1240 1250 1260
 CGCCACTCTGGTGGCTGGGCCGTCGGTGGCTGGCCGGTCCGGCT
 R H S W S R P L C S P L G R R R' A S
 P S V V V A A P V V A L G P A A E P G F 18/43
 T L G R G R C A R C A G A G G G P R L

 R S S S T V R T R T S P A P W C A Q P R
 G R H R R * G P E R H R L R G V H N R G
 V V I D G K D Q N V T G S V V C T T A A
 AGGTCTGTATCGACGGTAAGGACCAGAACGTCACCGGCTCCGTGGTGCACAACCGGG
 1270 1280 1290 1300 1310 1320
 TCCAGCAGTAGCTGCCATTCTGGTCTTGCAAGTGGCCGAGGCACACGTTGGGCC
 P R * R R Y P G S R * R S R P T C' L R P
 T T M S P L S W F T V P E T T H V V A A
 D D D V T L V L V D G A G H H A C G R G

FIGURE 6 (CONT'D)

P A M S T S R S A G R R P A L P P C S P
 R Q C Q H R D R R G G D R H C R R A H R
 G N V N I A I G G A A T G I A V L T D
 CCGGCAATGTCAACATGGCATCGCGATTGCCGGCATTGCTCACCG
 1330 1340 1350 1360 1370 1380 1390 1400 1410 1420 1430 1440 1450
 GGCGTTACAGTTGCTAGCGCTAGCCGGCGCTGGCGTAAACGGGACGGAGTGGC
 R C H * C R S R R P P S R C Q R R A * R
 P L T L M A I P P A V P M A A T S V S
 A I D V D R D A P R R G A N G G H 'E G V^{19/43}
 T A T L R R * S P L G S V T S T A S R W
 R Q P S G G E V R W A R * R Q R R H A G
 G N P P E V K S V G L G N V N G V T L G
 ACGGCAACCCTCGGAGGTGAAGTCCGGCTGGCTAACGTCACGGCTGG
 1390 1400 1410 1420 1430 1440 1450
 TGCGGTTGGAGGCCTCCACTTCAGGCAGGCATTGCAGTTGCCAGTGGGACC
 R C G E P P S T R Q A R Y R * R R * A P
 P L G G S T F D T P S P L T L P T V S ' P
 A V R R L H L G N P E T V D V A D R Q S

FIGURE 6 (CONT'D)

D T R R A P D R V T P R Q P R T A A T T
 I H V G H R T G * R L G N Q G R Q P L Q
 Y T S G T G Q G N A S A T K D G S H Y K
 GATACACGTGGCACCGAACGGTAACGGCTCGGCAACCAAGGAACGGCAGCCACTACA
 1450 1460 1470 1480 1490 1500
 20/43
 CTATGTCAGCCCCGTGGCCTGTCCCATTGCGGAGCCGTTGGTCTGCCGGTGGATGT
 I C T P C R V P Y R R P L W P R C G ' S C
 Y V D P V P C P L A E A V L S P L W * L
 V R R A G S L T V G R C G L V A A V V L
 R S L G P L P G S T W P T R C H R * T S
 D H W D R Y R G R H G Q P D V T G E Q V
 I T G T A T G V D M A N P M S P V N K S
 AGATCACTGGGACCGCTACCGGGTGGACATGGCCAACCCGATGTCACCGGGTGAACAAAGT
 1510 1520 1530 1540 1550 1560
 TCTAGTGAACCCCTGGCGATGGCCCCAGCTGTACAGTGGCCACTTGTCA
 S * Q S R * R P R C P W G S T V P S C T
 I V P V A V P T S M A L G I D G T F L D
 D S P G S G P D V H G V R H * R H V L R

FIGURE 6 (CONT'D)

R S K S R * P V P N L K R V D A G C E Q
 V R N R G D L F L T * S V S M R A V N S
 F E I E V T C S] * P K A C R C G L * T A
 CGTTCGAAATCGAGGTGACCTAACCTAAAGCGTGTGATGCCGGCTGTGAACAG
 1570 1580 1590 1600 1610 1620
 GCAAGCTTTAGCTCCACTGGACAAGGATTGGATTTCGCACAGCTACGCCCGACACTTGTC
 T R F R P S R N R V * L T D I R A T; F L
 N S I S T V Q E * G L A H R H P S H V A ^{21/43}
 E F D L H G T G L R F R T S A P Q S C R
 R V G A G Q S G L A R R F E R L P S V
 A S E P G S Q A * R G D D S S G C H P S
 R R S R A V R P S A A T I R A V A I R Q
 CGCGTGGAGCCGGCAGTCAGGCCCTAGCGGGCACGATTGAGCGGTGTCATCCGTC
 1630 1640 1650 1660 1670 1680
 GCGCAGCCTCGCCCCGTCAAGTCCGGATCGGCCAACGGTAGGCAG
 A D S G P L * A * R P S S E L P Q W G D
 R R L R A T L G L A A V I R A T A M R *
 T P A P C D P R A R R W S R N G D T L

FIGURE 6 (CONT'D)

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K	W	Q	P	H	R	K	L	G	I	S	G	*	A	T	H'	G	D	R	S
S	G	N	R	T	A	N	S	V	Y	P	G	E	L	L	T	V	I	V	P
V	A	T	A	P	Q	T	R	Y	I	R	V	S	Y	S	R	*	S	F	F
AAGTGGCAACCGCACCGAAACTCGGTATATCCGGGTGAGCTACTCACGGTGATCGTTCC																			
1690	1700	1710	1720	1730	1740														
TTCACCGTTGGCGTTTGAGCCATAAGGCCACTCGATGAGTGGCCACTAGCAAAGG																			
L	P	L	R	V	A	F	E	T	Y	G	P	S	S	V	T	I	T	G	
T	A	V	A	G	C	V	R	Y	I	R	T	L	*	E	R	H	D	N	R
H	C	G	C	R	L	S	P	I	D	P	H	A	V	*	P	S	R	E	T
V	V	R	L	D	H	S	G	D	D	R	Q	A	E	P	G	A	T	G	L
L	C	A	L	T	T	A	E	T	I	A	R	P	S	P	V	L	P	A	W
C	A	P	*	P	Q	R	R	R	S	P	G	R	A	R	C	Y	R	L	G
GTTGTGGCCTTGACCACAGCGGAGACGATGCCAGGGAGCCGCTACCGGGCTTG																			
1750	1760	1770	1780	1790	1790	1790	1790	1790	1790	1790	1790	1790	1790	1790	1790	1790	1790	1790	1790
CAGACGGAACTGGTGTGGCTCTGCGATGGCCACGGATGGCCGAAC																			
N	H	A	K	V	V	A	S	V	I	A	L	G	L	G	T	S	G	A	Q
Q	A	G	Q	G	C	R	L	R	D	G	P	R	A	R	H	*	R	S	P
T	R	R	S	W	L	P	S	S	R	W	A	S	G	P	A	V	P	K	A

FIGURE 6 (CONT'D)

A G P * R I A A G E P L E N L G L Q R G
 R D R D V S P R A N R S K T S D C S A A
 G T V T Y R R G R T A R K P R T A A R P
 GCGGACCGTGTACGGTATGCCGGCGAACCGCTCGAAAACCTCGGACTGCAGGCCGGC
 1810 1820 1830 1840 1850 1860
 CGCCCTGGCACTGCATAAGCGGGCGCCCGCTTGCGGAGCTTTGGAGCCTGACGTCGCCGG
 R S R S T D G R A F R E F V E S Q P A A
 P V T V Y R R P R V A R F G R V A A R G
 P G H R I A A P S G S S F R P S C R P R G^{23/43}
 R N T R P I V D H L Q H D V R R P G A Q
 G I P G P L S I T C S T T C V G P V L K
 E Y P A H C R S P A A R R A S A R C S S
 CGGAATACCCGCCCATTTGTCGATCACCTGCAGCACGTCGCTGGCCGGTGGCTCAA
 1870 1880 1890 1900 1910 1920
 GCCTTATGGCCGGGTAACAGCTMGTGGACGCTGCTGCACGGCAGCCACGAGTT
 P I G P G N D I V Q L V V H T P G T S ' L
 S Y G A W Q R D G A A R R A D A R H E L
 F V R G M T S * R C C S T R R G P A * A

FIGURE 6 (CONT'D)

A R R H D R A G T G V H A G V G Q Q V G
 R V V T I V P G P V C T R A L A S R L V
 A S S R S C R D R C A R G R W P A G W S
 GCGCGTCGTACCGATCGTGCCTGGGAGCCGGTGTGCACGCCAGGGTTGGT
 1930 1940 1950 1960 1970 1980 24/43
 CGCGCAGCAGTAGCAGGCCACACGTCGGCCACCGGTCTCGTCCAAACCA
 R T V I T G P G T H V R A N A L L' N T
 A D D R D H R S R H A R P R Q G A P Q D
 R R * S R A P V P T C A P T P W C T P *

 H H L V Q P C R I T R D D H R F G G Q V
 T T W C N R A A S P G M T T G S G G R S
 P P G A T V P H H P G * P P V R G A G R
 CACCACCTGGTGCAACCGTGCATCACCCGGATGACCAAGGGTCTGGGGCAGGTC
 1990 2000 2010 2020 2030 2040
 GTGGTGGACCACTGGCACGGCGTAGTGGCCCTACTGGTGGCCAAGCCCCGGTCCAG
 V V Q H L R A A D G P I V V P E P P L D
 G G P A V T G C * G P H G T R P A P R
 W R T C G H R M V R S S W R N P P C T P S

FIGURE 6 (CONT'D)

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E R P L V I W S G N M S V A D R V N R K	
S A H W * S G P A T * A S L T A S T A S	
A P T G D L V R Q H E R R * P R Q P Q A	
GAGCGCCCACTGGTGAATCTGGCTGAGCGTCAGCCGGCAAG	
2050 2060 2070 2080 2090 2100	
CTCGGGGTGACCTAGACCAGGGCTTACTCCAGGGACTGGCAGTTGGCGTTC	
L A W Q H D P G A V H A D S V A D V A L	
A G V P S R T R C C S R R Q G R * G C A	
R G S T I Q D P L M L T A S R T L R L G	
P R H V H R S A F Q R P P R V Q T R Q Q	
R D M S T G P R S S G R P E S R R A S S	
A T C P P V R V P A A A P S P D A P A A	
CCGGACATGTCCACCGGTTCCAGGGCCGGCCAGTGCCAGA	
2110 2120 2130 2140 2150 2160	
GGCGCTGTACAGGTGGCCAGGGCAAAGGTGGGGCTCAGGTCAGGGCTGGCTCGCGGGTC	
R S M D V P G R E L P R G S D L R A L L	
A V H G G T R T G A A G L G S A G A A	
R C T W R D A N W R G G R T W V R W C C	

FIGURE 6 (CONT'D)

Q Q V L D E T C Y P L G L R C H P A H R
 S R S S T R R V I R S V S D A T R L I A
 A G P R R D V L S A R S P M P P G S S R
 CAGCAGGTCCGACGGAGACGTATTCCGGATGCCACCCGGCTCATCGC
 2170 2180 2190 2200 2210 2220
 GTCGTCCAGGAGCTGCTCTGCACAATAGGGAGCCAGGGCTACGGTGGGGCGAGTAGCGC
 L L D E V L R T I R E T E S A V R S M A
 A P G R R S T N D A R D G I G G P E D R
 C T R S S V H * G S P R R H W G A * R T^{26/43}
 V C D G L G I V P Y P L R Q F R V T T D
 C A T A S G S S P I R C V N S V * P R I
 V R R P R D R P L S A A S I P C N H G S
 GTGTGCGACGGCCTCGGGATCGTCCCTATCCGCTGCGTCAATTCCGTGTAACCACGGAT
 2230 2240 2250 2260 2270 2280
 CACACGGCTGGAGCCCTAGCAGGGATAAGGCACATTGGTGCGCTA
 H A V A E P D D G I R Q T L E T Y G R I
 T R R G R S R G R D A A D I G H L W P D
 H S P R P I T G * G S R * N R T V V S R

FIGURE 6 (CONT'D)

R R K G S S Q F M T G I G N E L A H T T G
 A A R G V R S S * L A S A T N W R T R V
 P Q G E F A V H D W H R Q R T G A H G F
 CGCCGGAAAGGGAGTTCCGAGTTCATGACTGGCATCGGCCAACGGA
 2290 2300 2310 2320 2330 2340 27/43
 GCGGC GT TCCCCTCAAGCGTCAAGTACTGACCGTAGCCGTTGACCGGTGCCCCA
 A A L P T R L E H S A D A V F Q R V R T
 G C P S N A T * S Q C R C R V P A C P N
 R L P L E C N M V P M P L S S A C V P K
 F T G L P R R Q C G S D V V E H P V E R
 S L A C R A D S A A M W S S I R L S A
 H W P A A P T V R Q R C G R A S G * A P
 TTCACTGGCCTGCCGGACAGTGGCGATGTGGTCTGAGGCATCCGGTTGAGCGC
 2350 2360 2370 2380 2390 2400
 AAGTGACGGACGGCGGGCTGTCA CGCCGTACACCA GCTCGCTAGGCCAACTCGCG
 E S A Q R A S L A A A I H D L M R N L A
 * Q G A A G V T R C R H P R A D P Q A G
 V P R G R R C H P L S T T S C G T S R R

FIGURE 6 (CONT'D)

R P E L P H L G G V C V R F G H P D R
 D P S C P T S V E G F A S G T R T G
 T R V A P P R W R G L R Q V R A P G P V
 CGACCGAGTTGCCACCTCGGTGGAGGGTTGGCTCAGGTTGGCACCCGGACCGG
 2410 2420 2430 2440 2450 2460
 GCTGGCTAACGGGGTGGAGCCACCTCCCCAACGGCAGTCCAAGCCCCGTGGGCC
 S G L Q G V E T S P N A D P E P V R V P
 V R T A G G R H L P K R * T R A G P' G T
 G S N G W R P P T Q T L N P C G S R Y^{28/43}
 * L D L A A I Q R * V D D F A R G L R D
 S L T S P R S N G R S T T S L A V C A T
 A * P R R D P T V G R R L R S R F A R P
 TAGCTGACCTGCCCGATCCAACGGTAGGTGACACTTCGCTCGGGTTGCCGAC
 2470 2480 2490 2500 2510 2520 2530
 ATCGAACTGGAGCGGGCTAGGGTGGCATCCAGCTGCCATGGCTGAAGCGAGCGCCAAACGGCGCTG
 L K V E G R D L P L D V V E S A T Q A V
 A Q G R R S G V T P R R S R E R N A R G
 S S R A A I W R Y T S S K A R P K R S R

FIGURE 6 (CONT'D)

R R N G A S A R L M M T I P A V V A A T
 A A T A P A P A * * R F R R S S R R R P
 P Q R R Q R P L D D D S G G R R G D Q
 CGCGAACGGCCAGCCCCGCTGATGACGATTCCGGGTCTCGCGGCCGACC
 2530 2540 2550 2560 2570 2580
 GCGGCAGTGGCCGGTGGGGCGAACTACTACTGCTAAGGCCAGCAGGCCGCTGG
 A A V A G A G A Q H R N R D D R R G
 G C R R W R G S S S S E P P R R P S W
 R L P A L A R K I I V I G A T T A A V L
 N A I T V T I P K M I S I C N I V A S T
 T Q S P * R F R K * S A S A T S W R R R
 R N H R D D S E N D Q H L Q H R G V D V
 AACGCAATCACCGTGAACGATTCCGAAATGATCAGCATCGAACATCGTGGCGTCGACG
 2590 2600 2610 2620 2630 2640
 TTGCGTTAGTGGCACTGCTAAGGGCTTACTAGTCGTTAGACGTTGCTAGCACCCGAGCTGC
 V C D G H R N R F H D A D A V D H R R R
 R L * R S S E S F S * C R C C R P T S T
 A I V T V I G F I I L M Q L M T A D V N
 L P I D R P V T M T S C P F R L G A A S
 C P S T G R * R * R R A R F G S E R P A
 A H R Q A G D D V V P V S A R S G Q H
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FIGURE 7

1 GAA TTC CAA CCC TCG GTG CAG ATC CAG GTC TAT CAG GGG GAG CGT GAG 48
Glu Phe Gln Pro Ser Val Gln Ile Gln Val Tyr Gln Gly Glu Arg Glu

49 ATC GCC GCG CAC AAC AAG TTG CTC GGG TCC TTC GAG CTG ACC GGC ATC 96
Ile Ala Ala His Asn Lys Leu Leu Gly Ser Phe Glu Leu Thr Gly Ile

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97 CCG CCG CGG CCG CGG GGG ATT CCG CAG ATC GAG GTC ACT TTC GAC ATC 144
Pro Pro Ala Pro Arg Gly Ile Pro Gln Ile Val Thr Phe Asp Ile

145 GAC GCC AAC GGC ATT GTG CAC GTC ACC GCC AAG GAC AAG GGC ACC GGC 192
Asp Ala Asn Gly Ile Val His Val Thr Ala Lys Asp Lys Gly Thr Gly

FIGURE 7 (CONT'D)

3 AAG GAG AAC ACG ATC CGA ATC CAG GAA GGC TCG GGC CTG TCC AAG GAA 240
Lys Glu Asn Thr Ile Arg Ile Glu Gly Ser GLY Leu Ser Lys Glu

11 GAC ATT GAC CGC ATG ATC AAG GAC GCC GAA GCG CAC GCC GAG GAG GAT 288
Asp Ile Asp Arg MET Ile Lys Asp Ala Glu Ala His Ala Glu Glu Asp

19 CGC AAG CGT CGC GAG GAG GCC GAT GTT CGT AAT CAA GCC GAG ACA TTG 336
Arg Lys Arg Arg Glu Glu Ala Asp Val Arg Asn Gln Ala Glu Thr Leu

7 GTC TAC CAG ACG GAG AAG TTC GTC AAA GAA CAG CGT GAG GCC GAG GGT 384
Val Tyr Gln Thr Glu Lys Phe Val Lys Glu Gln Arg Glu Ala Glu GLY

FIGURE 7 (CONT'D)

385 GGT TCG AAG GTA CCT GAA GAC ACG CTC AAC AAG GTT GAT GCC GCG GTG 432
 Gly Ser Lys Val Pro Glu Asp Thr Leu Asn Lys Val Asp Ala Ala Val

433 GCG GAA GCG GAA GGC GGC ACT TGG CCG ATC GGA TAT TTC GGC CAT CAA 480
 Ala Glu Ala Glu Gly Gly Thr Trp Arg Ile Gly Tyr Phe Gly His Gln^{32/43}

481 GTC GGC GAT GGA GAA GCT GGG CCA GGA GTC GCA GGC TCT GGG GCA AGC 528
 Val Gly Asp Gly Glu Ala Gly Pro Gly Val Ala Gly Ser Gly Ala Ser

529 GAT CTA CGA AGC AGC TCA GGC TGC ACA GGC CAC TGG CGC TGC CCA 576
 Asp Leu Arg Ser Ser Gly Cys Val Thr Gly His Trp Arg Cys Pro

577 CCC CGG CGG CGA GCC GGC CGG TGC CCA CCC CGG CTC GGC 615
 Pro Arg Arg Arg Ala Gly Arg Cys Pro Pro Arg Leu Gly

FIGURE 8

5' TCGAACGAGGGGGTGAACCCGGTTGGCACTCGGCTTCTGGACTCGGCATAGGGGAGGTGCTTAAG
 3' AGCTTGCTCCCCGCACTGGCCACGGCTGAGGCCACTACGGATCCAGGGTATCCTGGCTCACGATT
 10 20 30 40 50
 70 80 90 100 110 120
 130 140 150 160 170 180
 190 200 210 220 230 240
 250 260 270 280 290 300
 310 320 330 340 350 360
 370 380 390 400
 410 420 430 440
 450 460 470 480
 490 500 510 520
 530 540 550 560
 570 580 590 595
 600 610 620 630
 640 650 660 670
 680 690 700 710
 720 730 740 750
 760 770 780 790
 800 810 820 830
 840 850 860 870
 880 890 900 910
 920 930 940 950
 960 970 980 990
 1000 1010 1020 1030
 1040 1050 1060 1070
 1080 1090 1100 1110
 1120 1130 1140 1150
 1160 1170 1180 1190
 1200 1210 1220 1230
 1240 1250 1260 1270
 1280 1290 1300 1310
 1320 1330 1340 1350
 1360 1370 1380 1390
 1400 1410 1420 1430
 1440 1450 1460 1470
 1480 1490 1500 1510
 1520 1530 1540 1550
 1560 1570 1580 1590
 1595 1600 1610 1620
 1630 1640 1650 1660
 1670 1680 1690 1700
 1710 1720 1730 1740
 1750 1760 1770 1780
 1790 1800 1810 1820
 1830 1840 1850 1860
 1870 1880 1890 1900
 1910 1920 1930 1940
 1950 1960 1970 1980
 1990 2000 2010 2020
 2030 2040 2050 2060
 2070 2080 2090 2100
 2110 2120 2130 2140
 2150 2160 2170 2180
 2190 2200 2210 2220
 2230 2240 2250 2260
 2270 2280 2290 2300
 2310 2320 2330 2340
 2350 2360 2370 2380
 2390 2400 2410 2420
 2430 2440 2450 2460
 2470 2480 2490 2500
 2510 2520 2530 2540
 2550 2560 2570 2580
 2590 2595 2600 2610
 2620 2630 2640 2650
 2660 2670 2680 2690
 2695 2700 2710 2720
 2730 2740 2750 2760
 2770 2780 2790 2795
 2800 2810 2820 2830
 2840 2850 2860 2870
 2880 2890 2900 2910
 2920 2930 2940 2950
 2960 2970 2980 2990
 2995 3000 3010 3020
 3030 3040 3050 3060
 3070 3080 3090 3095
 3100 3110 3120 3130
 3140 3150 3160 3170
 3180 3190 3200 3210
 3220 3230 3240 3250
 3260 3270 3280 3290
 3295 3300 3310 3320
 3330 3340 3350 3360
 3365 3370 3380 3390
 3395 3400 3410 3420
 3430 3440 3450 3460
 3470 3480 3490 3495
 3500 3510 3520 3530
 3540 3550 3560 3570
 3580 3590 3595 3600
 3605 3610 3620 3630
 3640 3650 3660 3670
 3680 3690 3695 3700
 3705 3710 3720 3730
 3740 3750 3760 3770
 3780 3790 3795 3800
 3805 3810 3820 3830
 3840 3850 3860 3870
 3880 3890 3895 3900
 3905 3910 3920 3930
 3940 3950 3960 3970
 3980 3990 3995 4000
 4005 4010 4020 4030
 4040 4050 4060 4070
 4080 4090 4095 4100
 4105 4110 4120 4130
 4140 4150 4160 4170
 4180 4190 4195 4200
 4205 4210 4220 4230
 4240 4250 4260 4270
 4280 4290 4295 4300
 4305 4310 4320 4330
 4340 4350 4360 4370
 4380 4390 4395 4400
 4405 4410 4420 4430
 4440 4450 4460 4470
 4480 4490 4495 4500
 4505 4510 4520 4530
 4540 4550 4560 4570
 4580 4590 4595 4600
 4605 4610 4620 4630
 4640 4650 4660 4670
 4680 4690 4695 4700
 4705 4710 4720 4730
 4740 4750 4760 4770
 4780 4790 4795 4800
 4805 4810 4820 4830
 4840 4850 4860 4870
 4880 4890 4895 4900
 4905 4910 4920 4930
 4940 4950 4960 4970
 4980 4990 4995 5000
 5005 5010 5020 5030
 5040 5050 5060 5070
 5080 5090 5095 5100
 5105 5110 5120 5130
 5140 5150 5160 5170
 5180 5190 5195 5200
 5205 5210 5220 5230
 5240 5250 5260 5270
 5280 5290 5295 5300
 5305 5310 5320 5330
 5340 5350 5360 5370
 5380 5390 5395 5400
 5405 5410 5420 5430
 5440 5450 5460 5470
 5480 5490 5495 5500
 5505 5510 5520 5530
 5540 5550 5560 5570
 5580 5590 5595 5600
 5605 5610 5620 5630
 5640 5650 5660 5670
 5680 5690 5695 5700
 5705 5710 5720 5730
 5740 5750 5760 5770
 5780 5790 5795 5800
 5805 5810 5820 5830
 5840 5850 5860 5870
 5880 5890 5895 5900
 5905 5910 5920 5930
 5940 5950 5960 5970
 5980 5990 5995 6000
 6005 6010 6020 6030
 6040 6050 6060 6070
 6080 6090 6095 6100
 6105 6110 6120 6130
 6140 6150 6160 6170
 6180 6190 6195 6200
 6205 6210 6220 6230
 6240 6250 6260 6270
 6280 6290 6295 6300
 6305 6310 6320 6330
 6340 6350 6360 6370
 6380 6390 6395 6400
 6405 6410 6420 6430
 6440 6450 6460 6470
 6480 6490 6495 6500
 6505 6510 6520 6530
 6540 6550 6560 6570
 6580 6590 6595 6600
 6605 6610 6620 6630
 6640 6650 6660 6670
 6680 6690 6695 6700
 6705 6710 6720 6730
 6740 6750 6760 6770
 6780 6790 6795 6800
 6805 6810 6820 6830
 6840 6850 6860 6870
 6880 6890 6895 6900
 6905 6910 6920 6930
 6940 6950 6960 6970
 6980 6990 6995 7000
 7005 7010 7020 7030
 7040 7050 7060 7070
 7080 7090 7095 7100
 7105 7110 7120 7130
 7140 7150 7160 7170
 7180 7190 7195 7200
 7205 7210 7220 7230
 7240 7250 7260 7270
 7280 7290 7295 7300
 7305 7310 7320 7330
 7340 7350 7360 7370
 7380 7390 7395 7400
 7405 7410 7420 7430
 7440 7450 7460 7470
 7480 7490 7495 7500
 7505 7510 7520 7530
 7540 7550 7560 7570
 7580 7590 7595 7600
 7605 7610 7620 7630
 7640 7650 7660 7670
 7680 7690 7695 7700
 7705 7710 7720 7730
 7740 7750 7760 7770
 7780 7790 7795 7800
 7805 7810 7820 7830
 7840 7850 7860 7870
 7880 7890 7895 7900
 7905 7910 7920 7930
 7940 7950 7960 7970
 7980 7990 7995 8000
 8005 8010 8020 8030
 8040 8050 8060 8070
 8080 8090 8095 8100
 8105 8110 8120 8130
 8140 8150 8160 8170
 8180 8190 8195 8200
 8205 8210 8220 8230
 8240 8250 8260 8270
 8280 8290 8295 8300
 8305 8310 8320 8330
 8340 8350 8360 8370
 8380 8390 8395 8400
 8405 8410 8420 8430
 8440 8450 8460 8470
 8480 8490 8495 8500
 8505 8510 8520 8530
 8540 8550 8560 8570
 8580 8590 8595 8600
 8605 8610 8620 8630
 8640 8650 8660 8670
 8680 8690 8695 8700
 8705 8710 8720 8730
 8740 8750 8760 8770
 8780 8790 8795 8800
 8805 8810 8820 8830
 8840 8850 8860 8870
 8880 8890 8895 8900
 8905 8910 8920 8930
 8940 8950 8960 8970
 8980 8990 8995 9000
 9005 9010 9020 9030
 9040 9050 9060 9070
 9080 9090 9095 9100
 9105 9110 9120 9130
 9140 9150 9160 9170
 9180 9190 9195 9200
 9205 9210 9220 9230
 9240 9250 9260 9270
 9280 9290 9295 9300
 9305 9310 9320 9330
 9340 9350 9360 9370
 9380 9390 9395 9400
 9405 9410 9420 9430
 9440 9450 9460 9470
 9480 9490 9495 9500
 9505 9510 9520 9530
 9540 9550 9560 9570
 9580 9590 9595 9600
 9605 9610 9620 9630
 9640 9650 9660 9670
 9680 9690 9695 9700
 9705 9710 9720 9730
 9740 9750 9760 9770
 9780 9790 9795 9800
 9805 9810 9820 9830
 9840 9850 9860 9870
 9880 9890 9895 9900
 9905 9910 9920 9930
 9940 9950 9960 9970
 9980 9990 9995 10000

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FIGURE 8 (CONT'D)

K E I E L E D P Y E K I G A E L V K E V
 AGGAGATCGAGCTGGAGGATCCGTACGAGAAGATGGGCCGAGCTGGTCAAGAGGTAG
 TCCCTAGCTCGAACCTCCCTAGGCATGCTCTCTAGGCCATGACCAGTTCTCCATC
 430 440 450 460 470 480

A K K T D D V A G D G T T A T V L A Q
 CCAAGAACCGATGACGTGGCTGGGGTGAACGGCACACAGAACGGGACGGGAA
 GGTCTTCCTGGCTACTGCCAGGGGCACTGCCGTGGTGGTGGCACGGACGGGTCC
 490 500 510 520 530 540

A L V R E G L R N V A A G A N P L G L K
 CGTTGGTTCCGGAGGGCCCTGGCAACGTCGGCCGGCCAAACCCGCTCGGTCTAAC
 GCAACCAAGGGCTCCGGACGGGTTGCAAGGGGGTGGGGGGGGGGGGGGGGGG
 550 560 570 580 590 600

R G I E K A V E K V T E T L K G A K E
 GCGGCATCGAAAAGGGGGGGAGGAAGGTACCGAGACCCCTGGCTCAAGGGGCCAAGGAGG
 CGCGTAGCTTTCCGGCACCTCTCCAGTGGCTGGGACGGAGTCCCCGGGTTCCTCC
 610 620 630 640 650 660

V E T K E Q I A A T A I S A G D Q S I
 TCGAGACCAAGGACCCAGATTGGGGTACCCGAGGGCATTTGGGGGTGACCCAGTCAG
 AGCTCTGGTCTGGCTAACGGCCGCTGGCTGGCTAAAGGCCCACTGGTCAGGTAGC
 670 680 690 700 710 720

G D L I A E M D K V G N E G V I T V E
 GTGACCTGATCGCCGAGGGATGGACAAGGGGGCAACGGAGGGGICATACCCGGTGGAGG
 CACTGGACTAGGGCTCCGGTACCTGTCACCCGTTGGCTCCCCGAGTAGTGGCAGCTCC
 730 740 750 760 770 780

FIGURE 8 (CONT'D)

E S H T F G L Q L E L T E G M R F D K G
 AGTCCAACACCTTTGGGCTTGAGCTCGAGCTCACCGAGGGTATGGGGTTCAGCAAGGGCT
 TCAGGGTTGTGGAAACCCGACGGCTCGAGCTAACGCCAAGCTGGTCCCCAAGCTGGTCCCCA
 790 800 810 820 830 840

 Y I S G Y F V T D P E R Q E A V L E D P
 ACATCTCGGGTACTTCGAGCCGACCCGGAGGGTCAAGGAGGGCTGGAGGGACCCCT
 TGTAGAGCCCCATGAAGCACTGGACTGGCTGGGCTCCGAGTCCCTGGGAGCCTCCTGGGA
 850 860 870 880 890 900

 Y I L V S K V S T V K D L L P L L E
 ACATCTGCTGGTCAGCTCCAAAGGIGGCCACTGTCAAGGATCTGGCTGGCTGGAGA
 TGTAGGACGACCAAGTGGAGGTCCACAGGTGACAGTCTAGACGGACGGGACGGAGCTCT
 910 920 930 940 950 960

 K V I G A G K P L L I I A E D V E G E A
 AGGTCAICGGAGGGGGTAAGGCCGCTGCTGATCATGCCGAGGACGCTGGAGGGCC
 TCCAGTAGCCCTGGCCATTCCGGGACTAGTGGGGCTCTGGAGCTCCCCTGGC
 970 980 990 1000 1010 1020

 L S T L V V N K I R G T F K S V A V K A
 TGTCACCCCTGGTCGICAACAAAGATCCGGCACCTTAAGGCTGGAGATTCAGGCT
 ACAGGTGGACCAAGGAGTGGTGTCTAGGGCCGTTCCAGGCTTACGGGAACTACGG
 1030 1040 1050 1060 1070 1080

 P G F G D R R K A M L Q D M A I L T G G
 CCGGCTTGGGCAACGGGGCAAGGGCAATGGCTGGAGGATATGGCCATTCACGGGCTC
 GGGCAAGGGCTGGGGTTCCGGTACGGACCTTACGGGAACTACGGTAAAGGTGGC
 1090 1100 1110 1120 1130 1140

 Q V I S E E V G L T L E H A D L S L L G
 AGGIGATCAGCGAAGAGGGGCGGCTGACGGCTGGAGAACGGGACCTGICGGCTGGCTAGGCA
 TCCACTAGTCGCTTCAGGGGACTGGGACCTGGGACCTGGACAGGGACGGATCCGT
 1150 1160 1170 1180 1190 1200

FIGURE 8 (CONT'D)

K A R K V V T K D E T T I V E G A G D
 AGGCCGGCAAGGGTGGTCACCAAGGAGACCACCATCGTGGAGGGGGGTGACA
 TCCGGGGGTTCCAGCACCGTGGTCTGGCTCTGGTAGCAGCTCCCCGGCCACTGT
 1210 1230 1220 1240 1250 1260

 T D A I A G R V A Q I R Q E I E N S D S
 CCGACGGCCATGGGGGACGGAGTGGCCAGATCCGCCAGGAGATCGAGAACGGGACTCCC
 GGCTGGGGTAGGGGCTGGTCAACGGGCTTAAGGGCTCTAGCTTGTGGCTGAGGC
 1270 1280 1300 1290 1310 1320

 D Y D R E K L Q E R L A K L A G G V A V
 ACTACGACCCGTGAGAAGCTGCAGGGAGGGCTGGCCAAGCTGGGCGGATCGGGCTGA
 TGATGCTGGCACTCTTGACGTCCCTGGGGTACCCACAGGGCAACT
 1330 1340 1350 1360 1370 1380

 I K A G A A T E V E L K E R K H R I E D
 TCAAGGGGGTGGCGGCCACCGAGGTGAACTCAAGGAGGGCAAGCACCCTGGGTGCGGCTGA
 AGTTCCGGCAACGGGGTGGCTCCAGCTTGAGTCCCTGGGTAGCTCCACT
 1390 1400 1410 1420 1430 1440

 A V R N A K A V E E G I V A G G G V T
 CGGTTCGCAAATGGCAAGGCTGGGGCATCGTCGGGCTGGGGTGTGACGG
 GCCAAGCGTTACGGTTCGGGGCAGCTCCCTGGTAGCAGGGCACCCACACTGG
 1450 1460 1470 1480 1490 1500

 L L Q A A P T L D E L K L E G D E A T G
 TGTGGCAAGGGGGCCCGACCCCTGGACGGAGCTGAAGGCTCGAAGGGGAGGGGACCCGGG
 ACAACGTTGGGGGCTGGGACCTGCTCGACTTCGAGCTCCGCTGGCTGGCC
 1510 1520 1530 1540 1550 1560

 A N I V K V A L E A P L K Q I A F N S G
 CCAACATCGIAGGGTGGGGCTGGAGGGGGCTGAAGGAGATCGGGCTCAACTCCGGG
 GGTGTAGGACTTCCACGGGACCTCCGGGAAGTTGAGGGCTAGCGTCA
 1570 1580 1590 1600 1610 1620

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FIGURE 8 (CONT'D)

L E P G V V A E K V R N L P A G H G L N
TGGAGCCGGGGGCGGGGTGGGAGAAGGTGGCAACCTGGCTGGCCACGGACTGAACG
ACCTGGGGGGGACCAACGGGCTTTCCACGGGTTGGACGGGGCTGACTTGC
1830 1840 1850 1660 1670 1680

A Q T G V Y E D L L A A G V A D P V K V
CTCAGAACCGGGTGTCTACGAGGATCTGCTCGCTGGCTGGCTGACCCGGTCAGGTGA
GAGTCTGGGACAGATGCTCCTAGACGGGACGGGAAAGGACTGGCCAGTCCACT
1690 1700 1710 1720 1730 1740

T R S A L Q N A A S I A G L F L T T E A
CCCGTTGGGCCCTGCAGAAATGGGGGCTCCATCGGGGCTGTCTGACCCAGGGCG
GGCAAGGGGGGACGGTCTACGGGGAGGTAGCCGACAAGGACTGGTGCTCCGG
1750 1760 1770 1780 1790 1800

V V A D K P E K E K A S V P G G D M G
TCGTTGCCGACAAGCCCCAAAAGGAGAAAGGCTTCCGTTCGGCTGGGGGACATGGGTG
AGCAACGGGCTGGCTGGCCCTTCCGAAGGCAAGGGCACCCGGCTGTACCCAC
1810 1820 1830 1840 1850 1860

G M D F *
GCATGGATTCTGACCCCCGGAGAAGTGGCAGGGAGGGGGGCTTGTGGGGCC
CGTACCTAAAGACTGGGGGGGCTTCAGGGCTCTCAGGGAAACACCCCC
1870 1880 1890 1900 1910 1920

GGGCTCTGGTTGGGAGCTACGGTAACCGGAGAACACACGGCAGTCGTGTAGGCAACCTT
CCCAGGAGACCAACCCCTCGATGCCATGGCTCTTGTGTCAGGACATCCGGTGGAA
1930 1940 1950 1960 1970 1980

TGGCCCTGTGGGGAGTCGGGGGGGGCTCTGGGTGAGGAGGGGGATGGGTACGA
ACGGGGACACCCCCCTACCCCCGGGAGGGCACGTGGGGCTACCCATGCT
1990 2000 2010 2020 2030 2040

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FIGURE 8 (CONT'D)

FIGURE 8 (CONT'D)

2530 2540 2550 2560 2570 2580
 CCCCAAGCCCGAGTCGTGCCGAGTTGACGAAGCTGGGTAGCTGGGCCAGGCT
 GGCCTTGGGTCAAGCAACGACGGCTCAACTGGCTTCGAGGCCATGGACCAAGGTCCCCGA
 R L G S D H S V F S P Y S T G P S

 2590 2600 2610 2620 2630 2640
 TCTAAGGCCCGGGTTTGGCCCCGAGCCAGCAGCCGACTGCCGCTACCGGGCTTGGGT
 AGATTCGGGCAAAACGGGGCTCGGTGGCGGCGGATGGCCCCAAGGCCAA
 R L G P H A G S G A A S G P H P H

 2650 2660 2670 2680 2690 2700
 CCTGAGTCAGGCCAACAGGAGCACTGGGGGGGACGGGGCTGTTGGTCAG
 CGGACTCAGGTCCGGGGTGTCTCGTGACCCGGGGCACAAACCAGTC
 G S D L G V P A S A P V P T H T L

 2710 2720 2730 2740 2750 2760 2780
 GCGGAGTTGAGGACCGTTCGCCAGGGCTGGTGGAGACCCGGCTGGATCCGGAGGGGGA
 CGGGCTCAACTCCTGCAAGGGGTCCGGCACAAACCTCTGGGGGGCAACTAGGCTCCGGCT
 G S N L V H A L G H Q L G G T S G L A S

 2770 2780 2790 2800 2810 2820
 GGGGAGGATGCCGAAACTCAAAGGGCCATGGCTCATGGGGGGTGGGTAGCCGGTA
 CGGCTCCATGGGGCTTGAGTTGGGGGGACCGAGTACGGGGCATGGGGCT
 A L I G S S L A A T S M G G T A Y G A S

 2830 2840 2850 2860 2870 2880
 CCTGACCAAGGCCGCCCTCCGAGGCCAGCCGGCTTCCTAAAGGGGGCTTGCATCCCCGC
 CGACTGGTTCGGGGAGGGCTGGGAAAGGATTCCGGGGCAAAACGGTAGGGGGCG
 S V L A A E S G A A S G L A A H Q M G A

 2890 2900 2910 2920 2930 2940
 GTTCAGAAGCTGGTGTGAGGGCTGCCCTGGCTGGGGCTGGGAGGGCTCCGGGGCAACTAACGGGGCT
 CAAGGTCTTGGACCAACTCCGACCAACTGGAAAGGGAGGGGACGGGACTTGGGGGG
 H W F S T H I S G A S G L G A H I T G S

FIGURE 8 (CONT'D)

2960 2960 2970 2980 2990 3000
 GGTCCCCGATGCCGGCTGTTCAAGGGAGCCGAATTCCCCGATGCCGATGTTTCCGCTGCCGA
 CCAGGGCTACGGGACAAAGTCCTCGGGCTTAAGGGCTAACGGCTACAAGGGCACGGCT
 T G I C S H L S G S H I G I H G S G S
 3010 3020 3030 3040 3050 3060
 GTTGAATAAGCCGACCGTGCCTGGTGGTCACTGGGAAAGCCGATGTTGCCGCTAACCGA
 CAACTTATTCGGCTGGCAACGGGACGGCTCAAGGGCTAACGGCTAACGGCATGGCT
 H F L G V H G T G S H G F G I H G S G S
 3070 3080 3090 3100 3110 3120
 GTTGAAGCCGCCAAACCCATCTGGTGAATCACCCTGATGCCGATCCCAGATATTCCCGCT
 CAACTTGGGGGGCTTGGTAGACCACACTAGTGGCCACACTAGGGCTTGGCTATAAGGGCGA
 H F G G F M Q H D G T I G F G I H N G S
 3130 3140 3150 3160 3170 3180
 ACCGGGTGGCCGAAGCCGATATTCCCGATGCCGAGGGTGGCCGAGGGCCAGGGTGGCT
 TGGCCACAACGGCTTTCGGCTATAAGGGCAGGGCTCCAAACGGCTCCAAACGGGA
 G T H G F G I H N G D G L H G L H N G S
 3190 3200 3210 3220 3230 3240
 GCGGGTGTGGCTGCCGATGTTGCCGATGCCGGTGTGCCGATGCCGATGTTGTTGTT
 CGGCCACAACGGGACGGCTAACGGCCACGGCCACAAACGGCTAACAAACAAACAA
 G T H G S G I H G T G T H G S G I H N H H
 3260 3270 3280 3290 3300
 GCCGATGTTGTTGCCGATGTTGCCGATGTTGCCGATGTTGCCGGCTGCCGATGCCGA
 CGGCTACAACAACGGCTACAAACAAACGGCTAACAAACGGCTAACAGCTAACGGCT
 G I H H G I H H G I H G S G T H N G F
 3310 3320 3330 3340 3350 3360
 GCCCAGATIGATCTGGCCGGTCTTGGCGATGTCGATGCCGAGGGTCCCGAAGAACCTGCTG
 CGGGCTTAACTAGCCGGAAAGAACGGCTAACGGCTAACAGCTAACGGCTAACGGCT
 G I H N K R I H N I G I H R I V H N . H

FIGURE 8 (CONT'D)

3370 3380 3390 3400 3410 3420
 CCAGGGCCAGTTGCGACGGCATCGAAGTGGTAACCAGCCATGCCGC
 GGTCCCCGGTCAACACGGCTGGGGGTCTGGCTAGCTTCACCATGGGTAGGGCA
 W P A L Q A V A A S A D F H Y G A M A A
 3430 3440 3450 3460 3470 3480
 CACGTCCAATGCCAACATTGGCTCGTATGCCGCCCTCGACGTCCATGAGGCCGGAGGGT
 GTGCAGGTTACGGGTAAACGAGCATACGGGGAGCTGCAGGTACTGCCCTGGCAA
 V D L A W M Q E Y A A E V D H L A P A H
 3490 3500 3510 3520 3530 3540
 CTGCCCAAAACGAGTTGGTAGCTGGCAGCAGCTGCATCAAGGCCACGGTACCCAC
 GACGGGTTGGTCAAGCATCGAACGGTCTGACCTAGTCCCCTGGCTAACGGGATGGTG
 Q G F W N T A A L L Q M L G R H A V V
 3550 3560 3570 3580 3590 3600
 TGCCGGCTGCACGGGTGGCCAGGGCCCTCGAACCGGGCTGGCTGTGCCCCATGGCCCTG
 ACGGCCGACGGTCCCACGGGGTGGTGGGGAGCTGGGGCACGGACAACGGTACGGGAC
 A P Q V T A A L A E F A T A M A Q
 3610 3620 3630 3640 3650 3660
 TGGGGCGCTGGTTGGCTGGGGCTGGCTGAGCCAGGGCTAGCTACTGGGTTGG
 ACGGGGGGGAACAAAGGGGACGGGACGGGACGGACTCGGTGATCCATGACCCAAACG
 A A Q E A Q A A T S L W A L Y Q T A
 3670 3680 3690 3700 3710 3720
 GACGGCCATCATCGGGGGGGACGGGACGGGACGGGACTAGTCAGTTGGATGTT
 CTGGGGGTAGTAGGGGGGGCTGGGTGGCTGGGGTGGATCAGTCAGGCTAAC
 V A M M A A S P G L W A G S T L E S T
 3730 3740 3750 3760 3770 3780
 GACGGGAGCCAAAGGGGACGGCATTTGACGGGAGCTGGGGGGGGGGGGGGGG
 CTGG
 V S G L S A I S A L L E E A L E G W A T

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FIGURE 8 (CONT'D)

3790 3800 3810 3820 3830 3840
 GGGCGGAAATTAGGGGICCCGACCCCCAACATCAGTCCCCGAATTGATCTC
 CCGGGCTCGTTAATCGCCAGGGCTGGGCTGGGCTTAACAGAG
 A A I L P G S G P G A F M L A S N I E 3850 3860
 TGGCGGAAACCACGGAAAATGGGGCTTGTCAAGCCGATCCAACCTAACCTAACCTCAC
 ACCGGCCGTTGGTGGCTTACGGGGAAACAGTCGGCTAGGTTGAATTGACAGTCGGCTGGC
 P P L W A F H P S T L R D L K V T L S R
 3910 3920 3930 3940 3950 3960
 TTGCCGTGGCGGTATCGGCACCTCAATACCACTCATCCTTGGGTCAACCTTGGGAGCC
 AACGGCACGGCATAGCCGTGAAGTTATGGTGAGTAGAAACCCCAGTAGAAACCTCGGG
 Q R P P I P V E I G S M K P T M K P A G
 3970 3980 3990 4000 4010 4020
 CCTAGGAACCGCCAGCTTACCTAGTCCCCTGGGTAGGGGACTGGGGGGATGGAGC
 GGATCCCTGGGGTGGATGGATCAGGGCCATCCCCGGCTGACGGGGGGCTACGGTGG
 R P V A L K G L P Y P G V P R S A A
 4030 4040 4050 4060 4070 4080
 TGAGGGCTGGCACCTGGCCCCGTAATGTCGGCTGGTATGGCAAGGACCCGACGGGGGG
 ACTCCAGACGGTGGACGGGGCATTACAGGGGACCATACCCGTTGGCTGGGCGGGGG
 S P R G G A G Y H R Q P L C R R P G
 4090 4100 4110 4120 4130 4140
 AAGAGTIGCTCGGGACGGGTTCACCCGGTTGATCGAACATGTCGACGGAAACTCACCGAGG
 TTCTCAAGAGGGCTGGCAAGTGGGCAACTAGCTTGTACAGCTGGCTGGGGGG
 L T A G R R T *
 4150 4160 4170 4180 4190 4200
 CCCTCAACGACCAACTGGCTGGCTACCCGGACCCCCAGGGCAACAGGATTGGGTGGC
 CGGAGTGGCTGGTGGGGGACGAAGGGGGTGGGGTGGGGTGGGGTGGGGTGGGGTGGC

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FIGURE 8 (CONT'D)

TGGCTCTGGCACAGGGCCGGGTGGCAGGATAACAGGTGCCCATGTGGCCGGCTGGAAAG
 ACCAGAACCGCTGCGGGGGCAACGTCCTATAATGCCACGGGTACACGGGGCACCTTC
 4210 4220 4230 4240 4250 4260
 AGGTGTGGACCCGGGACGGTTGGGGCTTGGGITAGATCTGCCGGGCACGGACA
 TCCACACCTGGGGCTGCCAACCCACCTGGCGAAACCCAAATCTAGACGGGGCTGGCTGT
 4270 4280 4290 4300 4310 4320
 CCCGATAATGGACACCCGGTCCCCGAGGGATGIGGGGAAGGTACGGGACGGAAATTG
 GGCCTATAACCTGTTGGCAAGGGTCCATGGGGCTGGCTTAAG
 4330 4340 4350 4360 4370 4380
 31
 51